

# SEQUENCE LISTING



<110> Prayaga, Sudhirdas K  
Taupier Jr, Raymond J  
Bandaru, Raj

<120> NOVEL POLYPEPTIDES HOMOLOGOUS TO THYMOSIN, EPHRIN A  
RECEPTORS, AND FIBROMODULIN, AND POLYNUCLEOTIDES  
ENCODING SAME

<130> 15966-585 CIP2

<140> 09/973,424

<141> 2001-10-09

<150> 60/159,805

<151> 1999-10-15

<150> 60/159,992

<151> 1999-10-18

<150> 60/160,952

<151> 1999-10-22

<150> 09/689,486

<151> 2000-10-12

<150> 09/687,276

<151> 2000-10-13

<160> 84

<170> PatentIn Ver. 2.1

<210> 1

<211> 430

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (61) .. (234)

<400> 1

gccagcagga gtgccatggt gagaggcact ggcagggaat gctaggattg ttttaagaaa 60

atg gca gac aaa cca gac ata ggg gaa atc gcc agc ttc aat aag gcc	108
Met Ala Asp Lys Pro Asp Ile Gly Glu Ile Ala Ser Phe Asn Lys Ala	
1 5 10 15	

aag ctg aag aaa aca gag atg cag gag aac acc ctg ctg acc aaa gag	156
Lys Leu Lys Lys Thr Glu Met Gln Glu Asn Thr Leu Leu Thr Lys Glu	
20 25 30	

gcc att gag cag gag aag cgg gtg aaa ttt cct aag agc ctg gag gat	204
Ala Ile Glu Gln Glu Lys Arg Val Lys Phe Pro Lys Ser Leu Glu Asp	
35 40 45	

tcc cta ccc ctg tca tct tcg aga ccc cag tagtaatgtg gaggaagaat 254  
 Ser Leu Pro Leu Ser Ser Ser Arg Pro Gln  
 50 55

caccacaaga tggacacaag ccacaaactg tgacgtgaac ctgggcactc cgtgctgatg 314  
 ccaccagcct gaggggtccct atgggtccaa tcagactgcc aaattctctg gtttgccctg 374  
 ggatattata gaaaattatt tgcgtgaata atgaaaacac agtcatggc aaaaaa 430

<210> 2  
 <211> 58  
 <212> PRT  
 <213> Homo sapiens

<400> 2  
 Met Ala Asp Lys Pro Asp Ile Gly Glu Ile Ala Ser Phe Asn Lys Ala  
 1 5 10 15  
 Lys Leu Lys Lys Thr Glu Met Gln Glu Asn Thr Leu Leu Thr Lys Glu  
 20 25 30  
 Ala Ile Glu Gln Glu Lys Arg Val Lys Phe Pro Lys Ser Leu Glu Asp  
 35 40 45  
 Ser Leu Pro Leu Ser Ser Ser Arg Pro Gln  
 50 55

<210> 3  
 <211> 13  
 <212> PRT  
 <213> Homo sapiens

<400> 3  
 Lys Leu Lys Lys Thr Glu Asn Thr Gln Glu Glu Lys Asn  
 1 5 10

<210> 4  
 <211> 3018  
 <212> DNA  
 <213> Homo sapiens

<400> 4  
 atggcccccg cccggggccg cctgccccct gcgctctggg tcgtcacggc cgcggcgggcg 60  
 gcggccacct gcgtgtccgc ggcgcgcggc gaagtgaatt tgctggacac gtcgaccatc 120  
 cacgggggact ggggctggct cacgtatccg gctcatgggt gggactccat caacgagggtg 180  
 gacgagtcct tccagcccat ccacacgtac caggtttgca acgtcatgag cccaaccag 240  
 aacaactggc tgcgcacgag ctgggtcccc cgagacggcg cccggcgcggt ctatgctgag 300  
 atcaagttaa ccctgcgcga ctgcaacagc atgcctggtg tgctgggcac ctgcaaggag 360  
 accttcaacc tctactacct ggagtcggac cgcgacctgg gggccagcac acaagaaagc 420  
 cagttcctca aaatcgacac cattgcggcc gacgagagct tcacaggtgc cgaccttgggt 480  
 gtgcggcgctc tcaagctcaa cacggagggtg cgcagtgtgg gtcccctcag caagcgcggc 540

```

ttctacctgg ccttccagga cataggtgcc tgcctggcca tctctctctt ccgcatctac 600
tataagaagt gccctgccat ggtgcgcaat ctggctgcct tctcggaggc agtgacgggg 660
gccgactcgt cctcactggg ggaggtgagg ggccagtgcg tgcggcactc agaggagcgg 720
gacacaccca agatgtactg cagcgcggag ggcgagtggc tctgccccat cggcaaatgc 780
gtgtgcagtg ccggctacga ggagcggcgg gatgcctgtg tggcctgtga gctgggcttc 840
tacaagtcag cccctgggga ccagctgtgt gcccgctgcc ctccccacag ccactccgca 900
gctccagccg cccaagcctg ccactgtgac ctacgtact accgtgcagc cctggacccg 960
ccgtcctcag cctgcacccg gccaccctcg gcaccagtga acctgatctc cagtgtgaat 1020
gggacatcag tgactctgga gtgggcccct cccctggacc cagggtggccg cagtgcacatc 1080
acctacaatg ccgtgtgccg ccgtgcccc tgggactga gccgctgcga ggcatgtggg 1140
agcggcaccg gctttgtgcc ccagcagaca agcctggtgc aggcagcct gctggtggcc 1200
aacctgctgg cccacatgaa ctactccttc tggatcgagg ccgtcaatgg cgtgtccgac 1260
ctgagccccg agccccgcgg gccgctgtg gtcaacatca ccacgaacca ggagccccg 1320
tcccaggtgg tggatgatccg tcaagagcgg gcggggcaga ccagcgtctc gctgctgtgg 1380
caggagcccg agcagccgaa cggcatcatc ctggagtatg agatcaagta ctacgagaag 1440
gacaaggaga tgcagagcta ctccaccctc aaggccgtca ccaccagagc caccgtctcc 1500
ggcctcaagc cgggcacccg ctacgtgttc cagggtccgag cccgcacctc agcaggctgt 1560
ggcgccttca gccaggccat ggaggtggag accgggaaac cccggccccg ctatgcacac 1620
aggaccattg tctggatctg cctgacgctc atcacgggcc tgggtggtgct tctgctcctg 1680
ctcatctgca agaagaggca ctgtggctac agcaaggcct tccaggactc ggacgaggag 1740
aagatgcact atcagaatgg acaggcaccc ccacctgtct tctgcctct gcacacccc 1800
ccgggaaagc tcccagagcc ccagttctat gcggaacccc acacctacga ggagccaggc 1860
cgggcggggc gcagtttcac tcgggagatc gaggcctcta ggatccacat cgagaaaatc 1920
atcggctctg gagactccgg ggaagtctgc tacgggaggc tgcgggtgcc agggcagcgg 1980
gatgtgcccc tggccatcaa ggccctcaaa gccggctaca cggagagaca gaggcgggac 2040
ttcctgagcg aggcgtccat catggggcaa ttcgaccatc ccaacatcat ccgcctcgag 2100
ggtgtcgtca ccctggccg cctggcaatg attgtgactg agtacatgga gaacggctct 2160
ctggacacct tccctgaggac ccacgacggg cagttcacca tcatgcagct ggtgggcatg 2220
ctgagaggag tgggtgccgg catgcgctac ctctcagacc tgggctatgt ccaccgagac 2280
ctggccgccc gcaacgtcct ggttgacagc aacctggtct gcaagggtgc tgacttcggg 2340
ccttcacggg tgetggagga cgaccggat gctgcctaca ccaccaggg cggaagatc 2400
cccattccgt ggacggcccc agaggccatc gccttcgca ccttctcctc ggccagcgac 2460
gtgtggagct tcggcgtggt catgtgggag gtgctggcct atggggagcg gccctactgg 2520
aacatgacca accgggatgt gatcagctct gtggaggagg ggtaccgct gcccgaccc 2580
atgggtgcc cccacgcct gcaccagctc atgctcgact gttggcaca ggaccggcg 2640
cagcggcctc gcttctccca gattgtcagt gtcctcgat cgctcatccg cagccctgag 2700
agtctcaggg ccaccgccac agtcagcagg tccccacccc ctgccttcgt ccggagctgc 2760
tttgacctcc gagggggcag cgggtggcgg gggggcctca ccgtggggga ctggctggac 2820
tccatccgca tgggcccgtg ccgagaccac ttcgctgcgg gcggatactc ctctctgggc 2880
atggtgttac gcatgaacgc ccaggacgtg cgcgcctgg gcacaccct catgggccac 2940
cagaagaaga tcctgggcag catcagacc atgcggggcc agctgaccag caccagggg 3000
ccccgcggc acctctga 3018

```

```

<210> 5
<211> 992
<212> PRT
<213> Homo sapiens

```

```

<400> 5
Met Ala Pro Ala Arg Gly Arg Leu Pro Pro Ala Leu Trp Val Val Thr
  1             5             10             15

Ala Ala Ala Ala Ala Ala Thr Cys Val Ser Ala Ala Arg Gly Glu Val
  20             25             30

Asn Leu Leu Asp Thr Ser Thr Ile His Gly Asp Trp Gly Trp Leu Thr

```

35	40	45
Tyr Pro Ala His Gly Trp Asp Ser Ile Asn Glu Val Asp Glu Ser Phe 50 55 60		
Gln Pro Ile His Thr Tyr Gln Val Cys Asn Val Met Ser Pro Asn Gln 65 70 75 80		
Asn Asn Trp Leu Arg Thr Ser Trp Val Pro Arg Asp Gly Ala Arg Arg 85 90 95		
Val Tyr Ala Glu Ile Lys Phe Thr Leu Arg Asp Cys Asn Ser Met Pro 100 105 110		
Gly Val Leu Gly Thr Cys Lys Glu Thr Phe Asn Leu Tyr Tyr Leu Glu 115 120 125		
Ser Asp Arg Asp Leu Gly Ala Ser Thr Gln Glu Ser Gln Phe Leu Lys 130 135 140		
Ile Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr Gly Ala Asp Leu Gly 145 150 155 160		
Val Arg Arg Leu Lys Leu Asn Thr Glu Val Arg Ser Val Gly Pro Leu 165 170 175		
Ser Lys Arg Gly Phe Tyr Leu Ala Phe Gln Asp Ile Gly Ala Cys Leu 180 185 190		
Ala Ile Leu Ser Leu Arg Ile Tyr Tyr Lys Lys Cys Pro Ala Met Val 195 200 205		
Arg Asn Leu Ala Ala Phe Ser Glu Ala Val Thr Gly Ala Asp Ser Ser 210 215 220		
Ser Leu Val Glu Val Arg Gly Gln Cys Val Arg His Ser Glu Glu Arg 225 230 235 240		
Asp Thr Pro Lys Met Tyr Cys Ser Ala Glu Gly Glu Trp Leu Val Pro 245 250 255		
Ile Gly Lys Cys Val Cys Ser Ala Gly Tyr Glu Glu Arg Arg Asp Ala 260 265 270		
Cys Val Ala Cys Glu Leu Gly Phe Tyr Lys Ser Ala Pro Gly Asp Gln 275 280 285		
Leu Cys Ala Arg Cys Pro Pro His Ser His Ser Ala Ala Pro Ala Ala 290 295 300		
Gln Ala Cys His Cys Asp Leu Ser Tyr Tyr Arg Ala Ala Leu Asp Pro 305 310 315 320		
Pro Ser Ser Ala Cys Thr Arg Pro Pro Ser Ala Pro Val Asn Leu Ile 325 330 335		
Ser Ser Val Asn Gly Thr Ser Val Thr Leu Glu Trp Ala Pro Pro Leu		

340										345					350						
Asp	Pro	Gly	Gly	Arg	Ser	Asp	Ile	Thr	Tyr	Asn	Ala	Val	Cys	Arg	Arg						
		355					360					365									
Cys	Pro	Trp	Ala	Leu	Ser	Arg	Cys	Glu	Ala	Cys	Gly	Ser	Gly	Thr	Arg						
	370					375					380										
Phe	Val	Pro	Gln	Gln	Thr	Ser	Leu	Val	Gln	Ala	Ser	Leu	Leu	Val	Ala						
385					390					395					400						
Asn	Leu	Leu	Ala	His	Met	Asn	Tyr	Ser	Phe	Trp	Ile	Glu	Ala	Val	Asn						
				405					410					415							
Gly	Val	Ser	Asp	Leu	Ser	Pro	Glu	Pro	Arg	Arg	Ala	Ala	Val	Val	Asn						
			420					425					430								
Ile	Thr	Thr	Asn	Gln	Ala	Ala	Pro	Ser	Gln	Val	Val	Val	Ile	Arg	Gln						
		435					440						445								
Glu	Arg	Ala	Gly	Gln	Thr	Ser	Val	Ser	Leu	Leu	Trp	Gln	Glu	Pro	Glu						
	450					455					460										
Gln	Pro	Asn	Gly	Ile	Ile	Leu	Glu	Tyr	Glu	Ile	Lys	Tyr	Tyr	Glu	Lys						
465					470					475					480						
Asp	Lys	Glu	Met	Gln	Ser	Tyr	Ser	Thr	Leu	Lys	Ala	Val	Thr	Thr	Arg						
				485					490					495							
Ala	Thr	Val	Ser	Gly	Leu	Lys	Pro	Gly	Thr	Arg	Tyr	Val	Phe	Gln	Val						
			500					505					510								
Arg	Ala	Arg	Thr	Ser	Ala	Gly	Cys	Gly	Arg	Phe	Ser	Gln	Ala	Met	Glu						
		515					520					525									
Val	Glu	Thr	Gly	Lys	Pro	Arg	Pro	Arg	Tyr	Asp	Thr	Arg	Thr	Ile	Val						
	530					535					540										
Trp	Ile	Cys	Leu	Thr	Leu	Ile	Thr	Gly	Leu	Val	Val	Leu	Leu	Leu	Leu						
545					550					555					560						
Leu	Ile	Cys	Lys	Lys	Arg	His	Cys	Gly	Tyr	Ser	Lys	Ala	Phe	Gln	Asp						
				565					570					575							
Ser	Asp	Glu	Glu	Lys	Met	His	Tyr	Gln	Asn	Gly	Gln	Ala	Pro	Pro	Pro						
		580						585					590								
Val	Phe	Leu	Pro	Leu	His	His	Pro	Pro	Gly	Lys	Leu	Pro	Glu	Pro	Gln						
		595					600					605									
Phe	Tyr	Ala	Glu	Pro	His	Thr	Tyr	Glu	Glu	Pro	Gly	Arg	Ala	Gly	Arg						
	610					615					620										
Ser	Phe	Thr	Arg	Glu	Ile	Glu	Ala	Ser	Arg	Ile	His	Ile	Glu	Lys	Ile						
625					630					635					640						
Ile	Gly	Ser	Gly	Asp	Ser	Gly	Glu	Val	Cys	Tyr	Gly	Arg	Leu	Arg	Val						

645	650	655
Pro Gly Gln Arg Asp Val Pro Val Ala Ile Lys Ala Leu Lys Ala Gly		
660	665	670
Tyr Thr Glu Arg Gln Arg Arg Asp Phe Leu Ser Glu Ala Ser Ile Met		
675	680	685
Gly Gln Phe Asp His Pro Asn Ile Ile Arg Leu Glu Gly Val Val Thr		
690	695	700
Arg Gly Arg Leu Ala Met Ile Val Thr Glu Tyr Met Glu Asn Gly Ser		
705	710	715
Leu Asp Thr Phe Leu Arg Thr His Asp Gly Gln Phe Thr Ile Met Gln		
725	730	735
Leu Val Gly Met Leu Arg Gly Val Gly Ala Gly Met Arg Tyr Leu Ser		
740	745	750
Asp Leu Gly Tyr Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val		
755	760	765
Asp Ser Asn Leu Val Cys Lys Val Ser Asp Phe Gly Leu Ser Arg Val		
770	775	780
Leu Glu Asp Asp Pro Asp Ala Ala Tyr Thr Thr Thr Gly Gly Lys Ile		
785	790	795
Pro Ile Arg Trp Thr Ala Pro Glu Ala Ile Ala Phe Arg Thr Phe Ser		
805	810	815
Ser Ala Ser Asp Val Trp Ser Phe Gly Val Val Met Trp Glu Val Leu		
820	825	830
Ala Tyr Gly Glu Arg Pro Tyr Trp Asn Met Thr Asn Arg Asp Val Ile		
835	840	845
Ser Ser Val Glu Glu Gly Tyr Arg Leu Pro Ala Pro Met Gly Cys Pro		
850	855	860
His Ala Leu His Gln Leu Met Leu Asp Cys Trp His Lys Asp Arg Ala		
865	870	875
Gln Arg Pro Arg Phe Ser Gln Ile Val Ser Val Leu Asp Ala Leu Ile		
885	890	895
Arg Ser Pro Glu Ser Leu Arg Ala Thr Ala Thr Val Ser Arg Cys Pro		
900	905	910
Pro Pro Ala Phe Val Arg Ser Cys Phe Asp Leu Arg Gly Gly Ser Gly		
915	920	925
Gly Gly Gly Gly Leu Thr Val Gly Asp Trp Leu Asp Ser Ile Arg Met		
930	935	940
Gly Arg Tyr Arg Asp His Phe Ala Ala Gly Gly Tyr Ser Ser Leu Gly		

945		950		955		960
Met Val Leu Arg	Met Asn Ala Gln Asp Val Arg	Ala Leu Gly Ile Thr				
	965		970			975
Leu Met Gly His Gln Lys Lys Ile Leu Gly Ser Ile Gln Thr Met Arg						
	980		985			990

<210> 6  
 <211> 2025  
 <212> DNA  
 <213> Homo sapiens

<400> 6

atggtggtgg	cacacccac	cgccactgcc	accaccacgc	ccactgccac	tgtcacggcc	60
accgttgtga	tgaccacggc	caccatggac	ctgcgggact	ggctgttcct	ctgctacggg	120
ctcatcgct	tcctgacgga	ggcatcgac	agcaccacct	gcccccggt	gtgccgctgc	180
gacaacggct	tcactactg	caacgaccgg	ggactcacat	ccatccccgc	agatatccct	240
gatgatgcca	ccacctcta	cctgcagaac	aaccagatca	acaacgccgg	catccccag	300
gacctcaaga	ccaaggtaa	cgtgcaggtc	atctacctat	acgagaatga	cctggatgag	360
ttccccatca	acctgccccg	ctccctccgg	gagctgcacc	tgcaggacaa	caatgtgcgc	420
accattgcca	gggactcgct	ggcccgcatc	ccgctgctgg	agaagctgca	cctggatgac	480
aactccgtgt	ccaccgtcag	cattgaggag	gacgccttcg	ccgacagcaa	acagctcaag	540
ctgctcttcc	tgagccggaa	ccacctgagc	agcatccccct	cggggctgcc	gcacacgctg	600
gaggagctgc	ggctggatga	caaccgcata	tcacccatcc	cgtgcatagc	cttcaagggc	660
ctcaacagcc	tgcggcgccct	ggtgctggac	ggtaacctgc	tggccaacca	gcgcacgcgc	720
gacgacacct	tcagccgcct	acagaacctc	acagagctct	cgtggtgctg	caattcgctg	780
gccgcgccac	ccctcaacct	gccagcgccc	cacctgcaga	agctctacct	gcaggacaat	840
gccatcagcc	acatcccccta	caacacgctg	gccaaagatgc	gtgagctgga	gcggctggac	900
ctgtccaaca	acaacctgac	cacgctgccc	cgcggcctgt	tcgacgacct	ggggaacctg	960
gccagctgc	tgtcaggaa	caacccttgg	ttttgtggct	gcaacctcat	gtggctgcgg	1020
gactgggtga	aggcacgggc	ggccgtggtc	aacgtgcggg	gcctcatgtg	ccagggccct	1080
gagaaggtcc	ggggcatggc	catcaaggac	attaccagcg	agatggacga	gtgttttgag	1140
acggggccgc	agggcggcgt	ggccaatgcg	gctgccaaga	ccacggccag	caaccacgcc	1200
tctgccacca	cgcgccaggg	ttccctgttt	accctcaagg	ccaaaaggcc	agggtgcgc	1260
ctccccgact	ccaacattga	ctaccccatg	gccacgggtg	atggcgccaa	gacctggcc	1320
atccacgtga	aggccctgac	ggcagactcc	atccgcatca	cgtggaaggc	cacgctcccc	1380
gcctctcttt	tccggctcag	ttggctgcgc	ctgggccaca	gccagccgt	gggctccatc	1440
acggagacct	tgggtcaggg	ggacaagaca	gagtacctgc	tgacagccct	ggagcccaag	1500
tccacctaca	tcacttgcac	ggtcaccatg	gagaccagca	atgcctacgt	agctgatgag	1560
acaccctgtg	gtgccaaggc	agagacagcc	gacagctatg	gccctaccac	cacactcaac	1620
caggagcaga	acgctggccc	catggcgagc	ctgcccctgg	cgggcatcat	cggcggggca	1680
gtggctctgg	tcttctcttt	cctggctcctg	ggggccatct	gctggtacgt	gcaccaggct	1740
ggcgagctgc	tgaccgggga	gagggcctac	aaccggggca	gcaggaaaaa	ggatgactat	1800
atggagtccag	ggaccaagaa	ggataactcc	atcctggaaa	tccgcggccc	tgggtgcag	1860
atgctgccca	tcaaccgta	ccgcgcaaaa	gaggagtacg	tgggtccacac	tatcttcccc	1920
tccaacggca	gcagcctctg	caaggccaca	cacaccattg	gctacggcac	cacgcggggc	1980
taccgggacg	gcggcatccc	cgacatagac	tactcctaca	catga		2025

<210> 7  
 <211> 674  
 <212> PRT

<213> Homo sapiens

<400> 7

Met	Val	Val	Ala	His	Pro	Thr	Ala	Thr	Ala	Thr	Thr	Thr	Pro	Thr	Ala
1				5					10					15	
Thr	Val	Thr	Ala	Thr	Val	Val	Met	Thr	Thr	Ala	Thr	Met	Asp	Leu	Arg
			20					25					30		
Asp	Trp	Leu	Phe	Leu	Cys	Tyr	Gly	Leu	Ile	Ala	Phe	Leu	Thr	Glu	Val
		35					40					45			
Ile	Asp	Ser	Thr	Thr	Cys	Pro	Ser	Val	Cys	Arg	Cys	Asp	Asn	Gly	Phe
	50					55					60				
Ile	Tyr	Cys	Asn	Asp	Arg	Gly	Leu	Thr	Ser	Ile	Pro	Ala	Asp	Ile	Pro
65					70					75					80
Asp	Asp	Ala	Thr	Thr	Leu	Tyr	Leu	Gln	Asn	Asn	Gln	Ile	Asn	Asn	Ala
				85					90					95	
Gly	Ile	Pro	Gln	Asp	Leu	Lys	Thr	Lys	Val	Asn	Val	Gln	Val	Ile	Tyr
		100						105					110		
Leu	Tyr	Glu	Asn	Asp	Leu	Asp	Glu	Phe	Pro	Ile	Asn	Leu	Pro	Arg	Ser
	115						120					125			
Leu	Arg	Glu	Leu	His	Leu	Gln	Asp	Asn	Asn	Val	Arg	Thr	Ile	Ala	Arg
	130					135					140				
Asp	Ser	Leu	Ala	Arg	Ile	Pro	Leu	Leu	Glu	Lys	Leu	His	Leu	Asp	Asp
145					150					155				160	
Asn	Ser	Val	Ser	Thr	Val	Ser	Ile	Glu	Glu	Asp	Ala	Phe	Ala	Asp	Ser
				165					170					175	
Lys	Gln	Leu	Lys	Leu	Leu	Phe	Leu	Ser	Arg	Asn	His	Leu	Ser	Ser	Ile
		180						185					190		
Pro	Ser	Gly	Leu	Pro	His	Thr	Leu	Glu	Glu	Leu	Arg	Leu	Asp	Asp	Asn
		195					200					205			
Arg	Ile	Ser	Thr	Ile	Pro	Leu	His	Ala	Phe	Lys	Gly	Leu	Asn	Ser	Leu
	210					215					220				
Arg	Arg	Leu	Val	Leu	Asp	Gly	Asn	Leu	Leu	Ala	Asn	Gln	Arg	Ile	Ala
225					230					235				240	
Asp	Asp	Thr	Phe	Ser	Arg	Leu	Gln	Asn	Leu	Thr	Glu	Leu	Ser	Leu	Val
				245					250					255	
Arg	Asn	Ser	Leu	Ala	Ala	Pro	Pro	Leu	Asn	Leu	Pro	Ser	Ala	His	Leu
		260						265					270		
Gln	Lys	Leu	Tyr	Leu	Gln	Asp	Asn	Ala	Ile	Ser	His	Ile	Pro	Tyr	Asn
	275						280					285			



Thr Leu Ala Lys Met Arg Glu Leu Glu Arg Leu Asp Leu Ser Asn Asn  
 290 295 300  
 Asn Leu Thr Thr Leu Pro Arg Gly Leu Phe Asp Asp Leu Gly Asn Leu  
 305 310 315 320  
 Ala Gln Leu Leu Leu Arg Asn Asn Pro Trp Phe Cys Gly Cys Asn Leu  
 325 330 335  
 Met Trp Leu Arg Asp Trp Val Lys Ala Arg Ala Ala Val Val Asn Val  
 340 345 350  
 Arg Gly Leu Met Cys Gln Gly Pro Glu Lys Val Arg Gly Met Ala Ile  
 355 360 365  
 Lys Asp Ile Thr Ser Glu Met Asp Glu Cys Phe Glu Thr Gly Pro Gln  
 370 375 380  
 Gly Gly Val Ala Asn Ala Ala Lys Thr Thr Ala Ser Asn His Ala  
 385 390 395 400  
 Ser Ala Thr Thr Pro Gln Gly Ser Leu Phe Thr Leu Lys Ala Lys Arg  
 405 410 415  
 Pro Gly Leu Arg Leu Pro Asp Ser Asn Ile Asp Tyr Pro Met Ala Thr  
 420 425 430  
 Gly Asp Gly Ala Lys Thr Leu Ala Ile His Val Lys Ala Leu Thr Ala  
 435 440 445  
 Asp Ser Ile Arg Ile Thr Trp Lys Ala Thr Leu Pro Ala Ser Ser Phe  
 450 455 460  
 Arg Leu Ser Trp Leu Arg Leu Gly His Ser Pro Ala Val Gly Ser Ile  
 465 470 475 480  
 Thr Glu Thr Leu Val Gln Gly Asp Lys Thr Glu Tyr Leu Leu Thr Ala  
 485 490 495  
 Leu Glu Pro Lys Ser Thr Tyr Ile Ile Cys Met Val Thr Met Glu Thr  
 500 505 510  
 Ser Asn Ala Tyr Val Ala Asp Glu Thr Pro Val Cys Ala Lys Ala Glu  
 515 520 525  
 Thr Ala Asp Ser Tyr Gly Pro Thr Thr Thr Leu Asn Gln Glu Gln Asn  
 530 535 540  
 Ala Gly Pro Met Ala Ser Leu Pro Leu Ala Gly Ile Ile Gly Gly Ala  
 545 550 555 560  
 Val Ala Leu Val Phe Leu Phe Leu Val Leu Gly Ala Ile Cys Trp Tyr  
 565 570 575  
 Val His Gln Ala Gly Glu Leu Leu Thr Arg Glu Arg Ala Tyr Asn Arg  
 580 585 590

Gly Ser Arg Lys Lys Asp Asp Tyr Met Glu Ser Gly Thr Lys Lys Asp  
           595                                600                                605  
 Asn Ser Ile Leu Glu Ile Arg Gly Pro Gly Leu Gln Met Leu Pro Ile  
       610                                615                                620  
 Asn Pro Tyr Arg Ala Lys Glu Glu Tyr Val Val His Thr Ile Phe Pro  
 625                                630                                635                                640  
 Ser Asn Gly Ser Ser Leu Cys Lys Ala Thr His Thr Ile Gly Tyr Gly  
                                 645                                650                                655  
 Thr Thr Arg Gly Tyr Arg Asp Gly Gly Ile Pro Asp Ile Asp Tyr Ser  
       660                                665                                670  
 Tyr Thr

<210> 8  
 <211> 22  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Ag190 Forward  
           PCR Primer Sequence

<400> 8  
 tggaggaaga atcaccacaa ga 22

<210> 9  
 <211> 27  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Ag190 Probe  
           PCR Primer Sequence

<400> 9  
 caagccacaa actgtgacgt gaacctg 27

<210> 10  
 <211> 19  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Ag190 Reverse  
           PCR Primer Sequence

<400> 10  
 gtggcatcag cacggagtg 19

<210> 11  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: Ag087 Forward  
         PCR Primer Sequence  
  
 <400> 11  
 cgcagtttca ctcgggagat 20  
  
 <210> 12  
 <211> 31  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: Ag087 Probe  
         PCR Primer Sequence  
  
 <400> 12  
 cctctaggat ccacatcgag aaaatcatcg g 31  
  
 <210> 13  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: Ag087 Reverse  
         PCR Primer Sequence  
  
 <400> 13  
 agcagacttc cccggagtct 20  
  
 <210> 14  
 <211> 31  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: NOV2 Forward  
         PCR Primer Sequence  
  
 <400> 14  
 ggatccgcg cgcggcgaagt gaatttgctg g 31  
  
 <210> 15  
 <211> 29  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: NOV2 Reverse  
 PCR Primer Sequence

<400> 15  
 ctcgagggtc ctggtgtcat agcggggcc 29

<210> 16  
 <211> 18  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: NOV2 S1 PCR  
 Primer Sequence

<400> 16  
 tacctggagt cggaccgc 18

<210> 17  
 <211> 18  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: NOV2 S2 PCR  
 Primer Sequence

<400> 17  
 gcggtccgac tccaggta 18

<210> 18  
 <211> 19  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: NOV2 S3 PCR  
 Primer Sequence

<400> 18  
 cagtgcgtgc ggcactcag 19

<210> 19  
 <211> 19  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: NOV2 S4 PCR  
 Primer Sequence

<400> 19  
tgagtgccgc acgcactgg 19

<210> 20  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: NOV2 S5 PCR  
Primer Sequence

<400> 20  
ctggacccag gtggccgc 18

<210> 21  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: NOV2 S6 PCR  
Primer Sequence

<400> 21  
gcggccacct gggtccag 18

<210> 22  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: NOV2 S7 PCR  
Primer Sequence

<400> 22  
cccgagcagc cgaacggc 18

<210> 23  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: NOV2 S8 PCR  
Primer Sequence

<400> 23  
gccgttcggc tgctcggg 18

<210> 24

<211> 27  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: NOV3 Forward  
         PCR Primer Sequence  
  
 <400> 24  
 ggatccacca cctgcccctc ggtgtgc 27  
  
 <210> 25  
 <211> 35  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: NOV3 Reverse  
         PCR Primer Sequence  
  
 <400> 25  
 ctcgaggcca gcgttctgct cctggttgag tgtgg 35  
  
 <210> 26  
 <211> 18  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: NOV3 S1 PCR  
         Primer Sequence  
  
 <400> 26  
 cgcaccattg ccaggac 18  
  
 <210> 27  
 <211> 18  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Description of Artificial Sequence: NOV3 S2 PCR  
         Primer Sequence  
  
 <400> 27  
 gtccctggca atggtgcg 18  
  
 <210> 28  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>

<223> Description of Artificial Sequence: NOV3 S3 PCR  
Primer Sequence

<400> 28  
ctggtgcgca attcgctggc c 21

<210> 29  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: NOV3 S4 PCR  
Primer Sequence

<400> 29  
ggccagcgaa ttgcgcacca g 21

<210> 30  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: NOV3 S5 PCR  
Primer Sequence

<400> 30  
cacgcctctg ccaccacg 18

<210> 31  
<211> 18  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: NOV3 S6 PCR  
Primer Sequence

<400> 31  
cgtggtggca gaggcgtg 18

<210> 32  
<211> 30  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: pSec-V5 His  
Forward Oligonucleotide Primer Sequence

<400> 32  
ctcgtcctcg agggtaagcc tatccctaac 30

<210> 33  
<211> 31  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: pSec-V5 His  
Reverse Oligonucleotide Primer Sequence

<400> 33  
ctcgtcgggc ccctgatcag cgggtttaaa c 31

<210> 34  
<211> 40  
<212> PRT  
<213> Homo sapiens

<400> 34  
Met Ala Asp Lys Pro Asp Met Gly Glu Ile Ala Ser Phe Asp Lys Ala  
1 5 10 15  
Lys Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Thr Leu Pro Thr Lys  
20 25 30  
Glu Thr Ile Glu Gln Glu Lys Arg  
35 40

<210> 35  
<211> 10  
<212> PRT  
<213> Homo sapiens

<400> 35  
Lys Leu Lys Lys Thr Glu Thr Gln Glu Asn  
1 5 10

<210> 36  
<211> 38  
<212> PRT  
<213> Homo sapiens

<400> 36  
Ala Asp Lys Pro Asp Met Gly Glu Ile Ala Ser Phe Asp Lys Ala Lys  
1 5 10 15  
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Thr Leu Pro Thr Lys Glu  
20 25 30  
Thr Ile Glu Gln Glu Lys  
35



<210> 37  
 <211> 40  
 <212> PRT  
 <213> Bos taurus

<400> 37  
 Ala Asp Lys Pro Asp Leu Gly Glu Ile Asn Ser Phe Asp Lys Ala Lys  
   1                  5                  10                  15  
 Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Thr Leu Pro Thr Lys Glu  
                   20                  25                  30  
 Thr Ile Glu Gln Glu Lys Gln Ala  
                   35                  40

<210> 38  
 <211> 40  
 <212> PRT  
 <213> Sus scrofa

<400> 38  
 Ala Asp Lys Pro Asp Met Gly Glu Ile Asn Ser Phe Asp Lys Ala Lys  
   1                  5                  10                  15  
 Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Thr Leu Pro Thr Lys Glu  
                   20                  25                  30  
 Thr Ile Glu Gln Glu Lys Gln Ala  
                   35                  40

<210> 39  
 <211> 40  
 <212> PRT  
 <213> Homo sapiens

<400> 39  
 Ser Asp Lys Pro Asp Met Ala Glu Ile Glu Lys Phe Asp Lys Ser Lys  
   1                  5                  10                  15  
 Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys Glu  
                   20                  25                  30  
 Thr Ile Glu Gln Glu Lys Gln Ala  
                   35                  40

<210> 40  
 <211> 41  
 <212> PRT  
 <213> Mus musculus

<400> 40  
 Met Ser Asp Lys Pro Asp Met Ala Glu Ile Glu Lys Phe Asp Lys Ser  
   1                  5                  10                  15

Lys Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys  
 20 25 30

Glu Thr Ile Glu Gln Glu Lys Gln Ala  
 35 40

<210> 41

<211> 40

<212> PRT

<213> *Oryctolagus cuniculus*

<400> 41

Ala Asp Lys Pro Asp Met Ala Glu Ile Glu Lys Phe Asp Lys Ser Lys  
 1 5 10 15

Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys Glu  
 20 25 30

Thr Ile Glu Gln Glu Lys Gln Ala  
 35 40

<210> 42

<211> 39

<212> PRT

<213> *Xenopus laevis*

<400> 42

Ser Asp Lys Pro Asp Met Ala Glu Ile Glu Lys Phe Asp Lys Ala Lys  
 1 5 10 15

Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys Glu  
 20 25 30

Thr Ile Glu Gln Glu Lys Gln  
 35

<210> 43

<211> 40

<212> PRT

<213> *Homo sapiens*

<400> 43

Ser Asp Lys Pro Gly Met Ala Glu Ile Glu Lys Phe Asp Lys Ser Lys  
 1 5 10 15

Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Ser Ser Lys Glu  
 20 25 30

Thr Ile Glu Gln Glu Arg Gln Ala  
 35 40

<210> 44

<211> 40

<212> PRT  
<213> Oncorhynchus mykiss

<400> 44  
Ser Asp Lys Pro Asn Leu Glu Glu Val Ala Ser Phe Asp Lys Thr Lys  
1 5 10 15  
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Thr Lys Glu  
20 25 30  
Thr Ile Glu Gln Glu Lys Gln Ala  
35 40

<210> 45  
<211> 40  
<212> PRT  
<213> Oncorhynchus mykiss

<400> 45  
Ser Asp Lys Pro Asp Leu Ala Glu Val Ser Asn Phe Asp Lys Thr Lys  
1 5 10 15  
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Thr Lys Glu  
20 25 30  
Thr Ile Glu Gln Glu Lys Gln Ala  
35 40

<210> 46  
<211> 40  
<212> PRT  
<213> Lateolabrax japonicus

<400> 46  
Ser Asp Lys Pro Asp Ile Ser Glu Val Thr Ser Phe Asp Lys Thr Lys  
1 5 10 15  
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys Glu  
20 25 30  
Thr Ile Glu Gln Glu Lys Ala Ala  
35 40

<210> 47  
<211> 39  
<212> PRT  
<213> Rattus norvegicus

<400> 47  
Met Ser Asp Lys Pro Asp Leu Ser Glu Val Glu Thr Phe Asp Lys Ser  
1 5 10 15  
Lys Leu Lys Lys Thr Asn Thr Glu Glu Lys Asn Thr Leu Pro Ser Lys  
20 25 30

Glu Thr Ile Gln Gln Glu Lys  
35

<210> 48  
<211> 38  
<212> PRT  
<213> Homo sapiens

<400> 48  
Ser Asp Lys Pro Asp Leu Ser Glu Val Glu Lys Phe Asp Arg Ser Lys  
1 5 10 15  
Leu Lys Lys Thr Asn Thr Glu Glu Lys Asn Thr Leu Pro Ser Lys Glu  
20 25 30

Thr Ile Gln Gln Glu Lys  
35

<210> 49  
<211> 35  
<212> PRT  
<213> Drosophila melanogaster

<400> 49  
Ile Ala Gly Ile Glu Asn Phe Asp Ala Lys Lys Leu Lys His Thr Glu  
1 5 10 15  
Thr Asn Glu Lys Asn Val Leu Pro Thr Lys Glu Val Ile Glu Ala Glu  
20 25 30

Lys Gln Ala  
35

<210> 50  
<211> 31  
<212> PRT  
<213> Drosophila melanogaster

<400> 50  
Gly Ile Thr Ala Phe Asn Gln Asn Asn Leu Lys His Thr Glu Thr Asn  
1 5 10 15  
Glu Lys Asn Pro Leu Pro Asp Lys Glu Ala Ile Glu Gln Glu Lys  
20 25 30

<210> 51  
<211> 38  
<212> PRT  
<213> Homo sapiens

<400> 51  
Ala Asp Lys Pro Asp Met Gly Glu Ile Ala Ser Phe Asp Lys Ala Lys

1	5	10	15
Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Thr Leu Pro Thr Lys Glu	20	25	30
Thr Ile Glu Gln Glu Lys	35		
<210> 52			
<211> 991			
<212> PRT			
<213> Mus musculus			
<400> 52			
Met Ala Pro Ala Arg Ala Arg Leu Ser Pro Ala Leu Trp Val Val Thr	5	10	15
1			
Ala Ala Ala Ala Ala Thr Cys Val Ser Ala Gly Arg Gly Glu Val Asn	20	25	30
Leu Leu Asp Thr Ser Thr Ile His Gly Asp Trp Gly Trp Leu Thr Tyr	35	40	45
Pro Ala His Gly Trp Asp Ser Ile Asn Glu Val Asp Glu Ser Phe Arg	50	55	60
Pro Ile His Thr Tyr Gln Val Cys Asn Val Met Ser Pro Asn Gln Asn	65	70	75
Asn Trp Leu Arg Thr Asn Trp Val Pro Arg Asp Gly Ala Arg Arg Val	85	90	95
Tyr Ala Glu Ile Lys Phe Thr Leu Arg Asp Cys Asn Ser Ile Pro Gly	100	105	110
Val Leu Gly Thr Cys Lys Glu Thr Phe Asn Leu His Tyr Leu Glu Ser	115	120	125
Asp Arg Asp Leu Gly Ala Ser Thr Gln Glu Ser Gln Phe Leu Lys Ile	130	135	140
Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr Gly Ala Asp Leu Gly Val	145	150	155
Arg Arg Leu Lys Leu Asn Thr Glu Val Arg Gly Val Gly Pro Leu Ser	165	170	175
Lys Arg Gly Phe Tyr Leu Ala Phe Gln Asp Ile Gly Ala Cys Leu Ala	180	185	190
Ile Leu Ser Leu Arg Ile Tyr Tyr Lys Lys Cys Pro Ala Met Val Arg	195	200	205
Asn Leu Ala Ala Phe Ser Glu Ala Val Thr Gly Ala Asp Ser Ser Ser	210	215	220

Leu Val Glu Val Arg Gly Gln Cys Val Arg His Ser Glu Glu Arg Asp  
 225 230 235 240  
 Thr Pro Lys Met Tyr Cys Ser Ala Glu Gly Glu Trp Leu Val Pro Ile  
 245 250 255  
 Gly Lys Cys Val Cys Ser Ala Gly Tyr Glu Glu Arg Arg Asp Ala Cys  
 260 265 270  
 Met Ala Cys Glu Leu Gly Phe Tyr Lys Ser Ala Pro Gly Asp Gln Leu  
 275 280 285  
 Cys Ala Arg Cys Pro Pro His Ser His Ser Ala Thr Pro Ala Ala Gln  
 290 295 300  
 Thr Cys Arg Cys Asp Leu Ser Tyr Tyr Arg Ala Ala Leu Asp Pro Pro  
 305 310 315 320  
 Ser Ala Ala Cys Thr Arg Pro Pro Ser Ala Pro Val Asn Leu Ile Ser  
 325 330 335  
 Ser Val Asn Gly Thr Ser Val Thr Leu Glu Trp Ala Pro Pro Leu Asp  
 340 345 350  
 Pro Gly Gly Arg Ser Asp Ile Thr Tyr Asn Ala Val Cys Arg Arg Cys  
 355 360 365  
 Pro Trp Ala Leu Ser His Cys Glu Ala Cys Gly Ser Gly Thr Arg Phe  
 370 375 380  
 Val Pro Gln Gln Thr Ser Leu Ala Gln Ala Ser Leu Leu Val Ala Asn  
 385 390 395 400  
 Leu Leu Ala His Met Asn Tyr Ser Phe Trp Ile Glu Ala Val Asn Gly  
 405 410 415  
 Val Ser Asn Leu Ser Pro Glu Pro Arg Ser Ala Ala Val Val Asn Ile  
 420 425 430  
 Thr Thr Asn Gln Ala Ala Pro Ser Gln Val Val Val Ile Arg Gln Glu  
 435 440 445  
 Arg Ala Gly Gln Thr Ser Val Ser Leu Leu Trp Gln Glu Pro Glu Gln  
 450 455 460  
 Pro Asn Gly Ile Ile Leu Glu Tyr Glu Ile Lys Tyr Tyr Glu Lys Asp  
 465 470 475 480  
 Lys Glu Met Gln Ser Tyr Ser Thr Leu Lys Ala Val Thr Thr Arg Ala  
 485 490 495  
 Thr Val Ser Gly Leu Lys Pro Gly Thr Arg Tyr Val Phe Gln Val Arg  
 500 505 510  
 Ala Arg Thr Ser Ala Gly Cys Gly Arg Phe Ser Gln Ala Met Glu Val  
 515 520 525

Glu	Thr	Gly	Lys	Pro	Arg	Pro	Arg	Tyr	Asp	Thr	Arg	Thr	Ile	Val	Trp
530						535					540				
Ile	Cys	Leu	Thr	Leu	Ile	Thr	Gly	Leu	Val	Val	Leu	Leu	Leu	Leu	Leu
545					550					555					560
Ile	Cys	Lys	Lys	Arg	His	Cys	Gly	Tyr	Ser	Lys	Ala	Phe	Gln	Asp	Ser
				565					570					575	
Asp	Glu	Glu	Lys	Met	His	Tyr	Gln	Asn	Gly	Gln	Ala	Pro	Pro	Pro	Val
			580					585					590		
Phe	Leu	Pro	Leu	Asn	His	Pro	Pro	Gly	Lys	Phe	Pro	Glu	Thr	Gln	Phe
		595					600					605			
Ser	Ala	Glu	Pro	His	Thr	Tyr	Glu	Glu	Pro	Gly	Arg	Ala	Gly	Arg	Ser
610						615					620				
Phe	Thr	Arg	Glu	Ile	Glu	Ala	Ser	Arg	Ile	His	Ile	Glu	Lys	Ile	Ile
625					630					635					640
Gly	Ser	Gly	Glu	Ser	Gly	Glu	Val	Cys	Tyr	Gly	Arg	Leu	Gln	Val	Pro
				645					650					655	
Gly	Gln	Arg	Asp	Val	Pro	Val	Ala	Ile	Lys	Ala	Leu	Lys	Ala	Gly	Tyr
			660					665					670		
Thr	Glu	Arg	Gln	Arg	Gln	Asp	Phe	Leu	Ser	Glu	Ala	Ala	Ile	Met	Gly
		675					680					685			
Gln	Phe	Asp	His	Pro	Asn	Ile	Ile	Arg	Leu	Glu	Gly	Val	Val	Thr	Arg
690					695						700				
Gly	Arg	Leu	Ala	Met	Ile	Val	Thr	Glu	Tyr	Met	Glu	Asn	Gly	Ser	Leu
705					710					715					720
Asp	Ala	Phe	Leu	Arg	Thr	His	Asp	Gly	Gln	Phe	Thr	Ile	Val	Gln	Leu
				725					730					735	
Val	Gly	Met	Leu	Arg	Gly	Val	Gly	Ala	Gly	Met	Arg	Tyr	Leu	Ser	Asp
			740					745					750		
Leu	Gly	Tyr	Ile	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Val	Leu	Val	Asp
		755					760					765			
Gly	Arg	Leu	Val	Cys	Lys	Val	Ser	Asp	Phe	Gly	Leu	Ser	Arg	Ala	Leu
770						775					780				
Glu	Asp	Asp	Pro	Glu	Ala	Ala	Tyr	Thr	Thr	Ala	Gly	Gly	Lys	Ile	Pro
785					790					795					800
Ile	Arg	Trp	Thr	Ala	Pro	Glu	Ala	Ile	Ala	Phe	Arg	Thr	Phe	Ser	Ser
				805					810					815	
Ala	Ser	Asp	Val	Trp	Ser	Phe	Gly	Val	Val	Met	Trp	Glu	Val	Leu	Ala
			820					825					830		

Tyr Gly Glu Arg Pro Tyr Trp Asn Met Thr Asn Gln Asp Val Ile Ser  
 835 840 845  
 Ser Val Glu Glu Gly Tyr Arg Leu Pro Ala Pro Met Gly Cys Pro Arg  
 850 855 860  
 Ala Leu His Gln Leu Met Leu Asp Cys Trp His Lys Asp Arg Ala Gln  
 865 870 875 880  
 Arg Pro Arg Phe Ala His Val Val Ser Val Leu Asp Ala Leu Val His  
 885 890 895  
 Ser Pro Glu Ser Leu Arg Ala Thr Ala Thr Val Ser Arg Cys Pro Pro  
 900 905 910  
 Pro Ala Phe Ala Arg Ser Cys Phe Asp Leu Arg Ala Gly Gly Ser Gly  
 915 920 925  
 Asn Gly Asp Leu Thr Val Gly Asp Trp Leu Asp Ser Ile Arg Met Gly  
 930 935 940  
 Arg Tyr Arg Asp His Phe Ala Ala Gly Gly Tyr Ser Ser Leu Gly Met  
 945 950 955 960  
 Val Leu Arg Met Asn Ala Gln Asp Val Arg Ala Leu Gly Ile Thr Leu  
 965 970 975  
 Met Gly His Gln Lys Lys Ile Leu Gly Ser Ile Gln Thr Met Arg  
 980 985 990

<210> 53  
 <211> 992  
 <212> PRT  
 <213> Homo sapiens

<400> 53  
 Met Ala Pro Ala Arg Gly Arg Leu Pro Pro Ala Leu Trp Val Val Thr  
 1 5 10 15  
 Ala Ala Ala Ala Ala Ala Thr Cys Val Ser Ala Ala Arg Gly Glu Val  
 20 25 30  
 Asn Leu Leu Asp Thr Ser Thr Ile His Gly Asp Trp Gly Trp Leu Thr  
 35 40 45  
 Tyr Pro Ala His Gly Trp Asp Ser Ile Asn Glu Val Asp Glu Ser Phe  
 50 55 60  
 Gln Pro Ile His Thr Tyr Gln Val Cys Asn Val Met Ser Pro Asn Gln  
 65 70 75 80  
 Asn Asn Trp Leu Arg Thr Ser Trp Val Pro Arg Asp Gly Ala Arg Arg  
 85 90 95  
 Val Tyr Ala Glu Ile Lys Phe Thr Leu Arg Asp Cys Asn Ser Met Pro  
 100 105 110



Gly Val Leu Gly Thr Cys Lys Glu Thr Phe Asn Leu Tyr Tyr Leu Glu  
 115 120 125  
 Ser Asp Arg Asp Leu Gly Ala Ser Thr Gln Glu Ser Gln Phe Leu Lys  
 130 135 140  
 Ile Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr Gly Ala Asp Leu Gly  
 145 150 155 160  
 Val Arg Arg Leu Lys Leu Asn Thr Glu Val Arg Ser Val Gly Pro Leu  
 165 170 175  
 Ser Lys Arg Gly Phe Tyr Leu Ala Phe Gln Asp Ile Gly Ala Cys Leu  
 180 185 190  
 Ala Ile Leu Ser Leu Arg Ile Tyr Tyr Lys Lys Cys Pro Ala Met Val  
 195 200 205  
 Arg Asn Leu Ala Ala Phe Ser Glu Ala Val Thr Gly Ala Asp Ser Ser  
 210 215 220  
 Ser Leu Val Glu Val Arg Gly Gln Cys Val Arg His Ser Glu Glu Arg  
 225 230 235 240  
 Asp Thr Pro Lys Met Tyr Cys Ser Ala Glu Gly Glu Trp Leu Val Pro  
 245 250 255  
 Ile Gly Lys Cys Val Cys Ser Ala Gly Tyr Glu Glu Arg Arg Asp Ala  
 260 265 270  
 Cys Val Ala Cys Glu Leu Gly Phe Tyr Lys Ser Ala Pro Gly Asp Gln  
 275 280 285  
 Leu Cys Ala Arg Cys Pro Pro His Ser His Ser Ala Ala Pro Ala Ala  
 290 295 300  
 Gln Ala Cys His Cys Asp Leu Ser Tyr Tyr Arg Ala Ala Leu Asp Pro  
 305 310 315 320  
 Pro Ser Ser Ala Cys Thr Arg Pro Pro Ser Ala Pro Val Asn Leu Ile  
 325 330 335  
 Ser Ser Val Asn Gly Thr Ser Val Thr Leu Glu Trp Ala Pro Pro Leu  
 340 345 350  
 Asp Pro Gly Gly Arg Ser Asp Ile Thr Tyr Asn Ala Val Cys Arg Arg  
 355 360 365  
 Cys Pro Trp Ala Leu Ser Arg Cys Glu Ala Cys Gly Ser Gly Thr Arg  
 370 375 380  
 Phe Val Pro Gln Gln Thr Ser Leu Val Gln Ala Ser Leu Leu Val Ala  
 385 390 395 400  
 Asn Leu Leu Ala His Met Asn Tyr Ser Phe Trp Ile Glu Ala Val Asn  
 405 410 415



Leu	Asp	Thr	Phe	Leu	Arg	Thr	His	Asp	Gly	Gln	Phe	Thr	Ile	Met	Gln
				725					730					735	
Leu	Val	Gly	Met	Leu	Arg	Gly	Val	Gly	Ala	Gly	Met	Arg	Tyr	Leu	Ser
			740					745					750		
Asp	Leu	Gly	Tyr	Val	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Val	Leu	Val
		755					760					765			
Asp	Ser	Asn	Leu	Val	Cys	Lys	Val	Ser	Asp	Phe	Gly	Leu	Ser	Arg	Val
	770					775					780				
Leu	Glu	Asp	Asp	Pro	Asp	Ala	Ala	Tyr	Thr	Thr	Thr	Gly	Gly	Lys	Ile
785					790					795					800
Pro	Ile	Arg	Trp	Thr	Ala	Pro	Glu	Ala	Ile	Ala	Phe	Arg	Thr	Phe	Ser
				805					810					815	
Ser	Ala	Ser	Asp	Val	Trp	Ser	Phe	Gly	Val	Val	Met	Trp	Glu	Val	Leu
			820					825					830		
Ala	Tyr	Gly	Glu	Arg	Pro	Tyr	Trp	Asn	Met	Thr	Asn	Arg	Asp	Val	Ile
		835					840					845			
Ser	Ser	Val	Glu	Glu	Gly	Tyr	Arg	Leu	Pro	Ala	Pro	Met	Gly	Cys	Pro
	850					855					860				
His	Ala	Leu	His	Gln	Leu	Met	Leu	Asp	Cys	Trp	His	Lys	Asp	Arg	Ala
865					870					875					880
Gln	Arg	Pro	Arg	Phe	Ser	Gln	Ile	Val	Ser	Val	Leu	Asp	Ala	Leu	Ile
				885					890					895	
Arg	Ser	Pro	Glu	Ser	Leu	Arg	Ala	Thr	Ala	Thr	Val	Ser	Arg	Cys	Pro
			900					905					910		
Pro	Pro	Ala	Phe	Val	Arg	Ser	Cys	Phe	Asp	Leu	Arg	Gly	Gly	Ser	Gly
		915					920					925			
Gly	Gly	Gly	Gly	Leu	Thr	Val	Gly	Asp	Trp	Leu	Asp	Ser	Ile	Arg	Met
	930					935					940				
Gly	Arg	Tyr	Arg	Asp	His	Phe	Ala	Ala	Gly	Gly	Tyr	Ser	Ser	Leu	Gly
945					950					955					960
Met	Val	Leu	Arg	Met	Asn	Ala	Gln	Asp	Val	Arg	Ala	Leu	Gly	Ile	Thr
				965					970					975	
Leu	Met	Gly	His	Gln	Lys	Lys	Ile	Leu	Gly	Ser	Ile	Gln	Thr	Met	Arg
			980					985					990		

<211> 450  
 <212> PRT  
 <213> Mus musculus

<400> 54

```

Met Ala Pro Ala Arg Ala Arg Leu Ser Pro Ala Leu Trp Val Val Thr
  1              5              10              15

Ala Ala Ala Ala Ala Thr Cys Val Ser Ala Gly Arg Gly Glu Val Asn
      20              25              30

Leu Leu Asp Thr Ser Thr Ile His Gly Asp Trp Gly Trp Leu Thr Tyr
      35              40              45

Pro Ala His Gly Trp Asp Ser Ile Asn Glu Val Asp Glu Ser Phe Arg
      50              55              60

Pro Ile His Thr Tyr Gln Val Cys Asn Val Met Ser Pro Asn Gln Asn
      65              70              75              80

Asn Trp Leu Arg Thr Asn Trp Val Pro Arg Asp Gly Ala Arg Arg Val
      85              90              95

Tyr Ala Glu Ile Lys Phe Thr Leu Arg Asp Cys Asn Ser Ile Pro Gly
      100             105             110

Val Leu Gly Thr Cys Lys Glu Thr Phe Asn Leu His Tyr Leu Glu Ser
      115             120             125

Asp Arg Asp Leu Gly Ala Ser Thr Gln Glu Ser Gln Phe Leu Lys Ile
      130             135             140

Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr Gly Ala Asp Leu Gly Val
      145             150             155             160

Arg Arg Leu Lys Leu Asn Thr Glu Val Arg Gly Val Gly Pro Leu Ser
      165             170             175

Lys Arg Gly Phe Tyr Leu Ala Phe Gln Asp Ile Gly Ala Cys Leu Ala
      180             185             190

Ile Leu Ser Leu Arg Ile Tyr Tyr Lys Lys Cys Pro Ala Met Val Arg
      195             200             205

Asn Leu Ala Ala Phe Ser Glu Ala Val Thr Gly Ala Asp Ser Ser Ser
      210             215             220

Leu Val Glu Val Arg Gly Gln Cys Val Arg His Ser Glu Glu Arg Asp
      225             230             235             240

Thr Pro Lys Met Tyr Cys Ser Ala Glu Gly Glu Trp Leu Val Pro Ile
      245             250             255

Gly Lys Cys Val Cys Ser Ala Gly Tyr Glu Glu Arg Arg Asp Ala Cys
      260             265             270

Met Ala Cys Glu Leu Gly Phe Tyr Lys Ser Ala Pro Gly Asp Gln Leu

```

275					280					285					
Cys	Ala	Arg	Cys	Pro	Pro	His	Ser	His	Ser	Ala	Thr	Pro	Ala	Ala	Gln
	290					295					300				
Thr	Cys	Arg	Cys	Asp	Leu	Ser	Tyr	Tyr	Arg	Ala	Ala	Leu	Asp	Pro	Pro
305					310					315					320
Ser	Ala	Ala	Cys	Thr	Arg	Pro	Pro	Ser	Ala	Pro	Val	Asn	Leu	Ile	Ser
				325					330					335	
Ser	Val	Asn	Gly	Thr	Ser	Val	Thr	Leu	Glu	Trp	Ala	Pro	Pro	Leu	Asp
			340					345					350		
Pro	Gly	Gly	Arg	Ser	Asp	Ile	Thr	Tyr	Asn	Ala	Val	Cys	Arg	Arg	Cys
		355					360					365			
Pro	Trp	Ala	Leu	Ser	His	Cys	Glu	Ala	Cys	Gly	Ser	Gly	Thr	Arg	Phe
	370					375						380			
Val	Pro	Gln	Gln	Thr	Ser	Leu	Ala	Gln	Ala	Ser	Leu	Leu	Val	Ala	Asn
385					390					395					400
Leu	Leu	Ala	His	Met	Asn	Tyr	Ser	Phe	Trp	Ile	Glu	Ala	Val	Asn	Gly
				405					410					415	
Val	Ser	Asn	Leu	Ser	Pro	Glu	Pro	Arg	Ser	Ala	Ala	Val	Val	Asn	Ile
			420					425					430		
Thr	Thr	Asn	Gln	Ala	Ala	Pro	Ser	Gln	Val	Val	Val	Ile	Arg	Gln	Glu
		435					440					445			
Arg	Ala														
	450														

<210> 55

<211> 480

<212> PRT

<213> Homo sapiens

<400> 55

Met	Arg	Gly	Ser	Gly	Pro	Arg	Gly	Ala	Gly	His	Arg	Arg	Pro	Pro	Ser
1				5					10					15	
Gly	Gly	Gly	Asp	Thr	Pro	Ile	Thr	Pro	Ala	Ser	Leu	Ala	Gly	Cys	Tyr
			20					25					30		
Ser	Ala	Pro	Arg	Arg	Ala	Pro	Leu	Trp	Thr	Cys	Leu	Leu	Leu	Cys	Ala
		35					40					45			
Ala	Leu	Arg	Thr	Leu	Leu	Ala	Ser	Pro	Ser	Asn	Glu	Val	Asn	Leu	Leu
	50					55					60				
Asp	Ser	Arg	Thr	Val	Met	Gly	Asp	Leu	Gly	Trp	Ile	Ala	Phe	Pro	Lys
65					70					75					80



Gly Arg Lys Asp Val Ser Tyr Tyr Ile Ala Cys Lys Lys Cys Asn Ser  
 385 390 395 400  
 His Ala Gly Val Cys Glu Glu Cys Gly Gly His Val Arg Tyr Leu Pro  
 405 410 415  
 Arg Gln Ser Gly Leu Lys Asn Thr Ser Val Met Met Val Asp Leu Leu  
 420 425 430  
 Ala His Thr Asn Tyr Thr Phe Glu Ile Glu Ala Val Asn Gly Val Ser  
 435 440 445  
 Asp Leu Ser Pro Gly Ala Arg Gln Tyr Val Ser Val Asn Val Thr Thr  
 450 455 460  
 Asn Gln Ala Ala Pro Ser Pro Val Thr Asn Val Lys Lys Gly Lys Ile  
 465 470 475 480

<210> 56  
 <211> 456  
 <212> PRT  
 <213> Gallus gallus

<400> 56  
 Met Gly Leu Arg Gly Gly Gly Gly Arg Ala Gly Gly Pro Ala Pro Gly  
 1 5 10 15  
 Trp Thr Cys Leu Leu Leu Cys Ala Ala Leu Arg Ser Leu Leu Ala Ser  
 20 25 30  
 Pro Gly Ser Glu Val Asn Leu Leu Asp Ser Arg Thr Val Met Gly Asp  
 35 40 45  
 Leu Gly Trp Ile Ala Tyr Pro Lys Asn Gly Trp Glu Glu Ile Gly Glu  
 50 55 60  
 Val Asp Glu Asn Tyr Ala Pro Ile His Thr Tyr Gln Val Cys Lys Val  
 65 70 75 80  
 Met Glu Gln Asn Gln Asn Asn Trp Leu Leu Thr Ser Trp Ile Ser Asn  
 85 90 95  
 Glu Gly Arg Pro Ala Ser Ser Phe Glu Leu Lys Phe Thr Leu Arg Asp  
 100 105 110  
 Cys Asn Ser Leu Pro Gly Gly Leu Gly Thr Cys Lys Glu Thr Phe Asn  
 115 120 125  
 Met Tyr Tyr Phe Glu Ser Asp Asp Glu Asp Gly Arg Asn Ile Arg Glu  
 130 135 140  
 Asn Gln Tyr Ile Lys Ile Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr  
 145 150 155 160

Glu Leu Asp Leu Gly Asp Arg Val Met Lys Leu Asn Thr Glu Val Arg  
 165 170 175  
 Asp Val Gly Pro Leu Thr Lys Lys Gly Phe Tyr Leu Ala Phe Gln Asp  
 180 185 190  
 Val Gly Ala Cys Ile Ala Leu Val Ser Val Arg Val Tyr Tyr Lys Lys  
 195 200 205  
 Cys Pro Ser Val Ile Arg Asn Leu Ala Arg Phe Pro Asp Thr Ile Thr  
 210 215 220  
 Gly Ala Asp Ser Ser Gln Leu Leu Glu Val Ser Gly Val Cys Val Asn  
 225 230 235 240  
 His Ser Val Thr Asp Glu Ala Pro Lys Met His Cys Ser Ala Glu Gly  
 245 250 255  
 Glu Trp Leu Val Pro Ile Gly Lys Cys Leu Cys Lys Ala Gly Tyr Glu  
 260 265 270  
 Glu Lys Asn Asn Thr Cys Gln Val Cys Arg Pro Gly Phe Phe Lys Ala  
 275 280 285  
 Ser Pro His Ser Pro Ser Cys Ser Lys Cys Pro Pro His Ser Tyr Thr  
 290 295 300  
 Leu Asp Glu Ala Ser Thr Ser Cys Leu Cys Glu Glu His Tyr Phe Arg  
 305 310 315 320  
 Arg Glu Ser Asp Pro Pro Thr Met Ala Cys Thr Arg Pro Pro Ser Ala  
 325 330 335  
 Pro Arg Ser Ala Ile Ser Asn Val Asn Glu Thr Ser Val Phe Leu Glu  
 340 345 350  
 Trp Ile Pro Pro Ala Asp Thr Gly Gly Arg Lys Asp Val Ser Tyr Tyr  
 355 360 365  
 Ile Ala Cys Lys Lys Cys Asn Ser His Ser Gly Leu Cys Glu Ala Cys  
 370 375 380  
 Gly Ser His Val Arg Tyr Leu Pro Gln Gln Thr Gly Leu Lys Asn Thr  
 385 390 395 400  
 Ser Val Met Met Val Asp Leu Leu Ala His Thr Asn Tyr Thr Phe Glu  
 405 410 415  
 Ile Glu Ala Val Asn Gly Val Ser Asp Gln Asn Pro Gly Ala Arg Gln  
 420 425 430  
 Phe Val Ser Val Asn Val Thr Thr Asn Gln Ala Ala Pro Ser Pro Val  
 435 440 445  
 Ser Ser Val Lys Lys Gly Lys Ile  
 450 455



<210> 57  
 <211> 649  
 <212> PRT  
 <213> Homo sapiens

<400> 57

```

Met Ile Ser Ala Ala Trp Ser Ile Phe Leu Ile Gly Thr Lys Ile Gly
  1               5               10               15

Leu Phe Leu Gln Val Ala Pro Leu Ser Val Met Ala Lys Ser Cys Pro
      20               25               30

Ser Val Cys Arg Cys Asp Ala Gly Phe Ile Tyr Cys Asn Asp Arg Phe
      35               40               45

Leu Thr Ser Ile Pro Thr Gly Ile Pro Glu Asp Ala Thr Thr Leu Tyr
  50               55               60

Leu Gln Asn Asn Gln Ile Asn Asn Ala Gly Ile Pro Ser Asp Leu Lys
  65               70               75               80

Asn Leu Leu Lys Val Glu Arg Ile Tyr Leu Tyr His Asn Ser Leu Asp
      85               90               95

Glu Phe Pro Thr Asn Leu Pro Lys Tyr Val Lys Glu Leu His Leu Gln
      100              105              110

Glu Asn Asn Ile Arg Thr Ile Thr Tyr Asp Ser Leu Ser Lys Ile Pro
      115              120              125

Tyr Leu Glu Glu Leu His Leu Asp Asp Asn Ser Val Ser Ala Val Ser
      130              135              140

Ile Glu Glu Gly Ala Phe Arg Asp Ser Asn Tyr Leu Arg Leu Leu Phe
      145              150              155              160

Leu Ser Arg Asn His Leu Ser Thr Ile Pro Trp Gly Leu Pro Arg Thr
      165              170              175

Ile Glu Glu Leu Arg Leu Asp Asp Asn Arg Ile Ser Thr Ile Ser Ser
      180              185              190

Pro Ser Leu Gln Gly Leu Thr Ser Leu Lys Arg Leu Val Leu Asp Gly
      195              200              205

Asn Leu Leu Asn Asn His Gly Leu Gly Asp Lys Val Phe Phe Asn Leu
      210              215              220

Val Asn Leu Thr Glu Leu Ser Leu Val Arg Asn Ser Leu Thr Ala Ala
      225              230              235              240

Pro Val Asn Leu Pro Gly Thr Asn Leu Arg Lys Leu Tyr Leu Gln Asp
      245              250              255

Asn His Ile Asn Arg Val Pro Pro Asn Ala Phe Ser Tyr Leu Arg Gln
  
```

260										265					270				
Leu	Tyr	Arg	Leu	Asp	Met	Ser	Asn	Asn	Asn	Leu	Ser	Asn	Leu	Pro	Gln				
		275						280				285							
Gly	Ile	Phe	Asp	Asp	Leu	Asp	Asn	Ile	Thr	Gln	Leu	Ile	Leu	Arg	Asn				
	290					295					300								
Asn	Pro	Trp	Tyr	Cys	Gly	Cys	Lys	Met	Lys	Trp	Val	Arg	Asp	Trp	Leu				
305					310					315					320				
Gln	Ser	Leu	Pro	Val	Lys	Val	Asn	Val	Arg	Gly	Leu	Met	Cys	Gln	Ala				
				325					330					335					
Pro	Glu	Lys	Val	Arg	Gly	Met	Ala	Ile	Lys	Asp	Leu	Asn	Ala	Glu	Leu				
			340					345					350						
Phe	Asp	Cys	Lys	Asp	Ser	Gly	Ile	Val	Ser	Thr	Ile	Gln	Ile	Thr	Thr				
	355					360						365							
Ala	Ile	Pro	Asn	Thr	Val	Tyr	Pro	Ala	Gln	Gly	Gln	Trp	Pro	Ala	Pro				
370					375						380								
Val	Thr	Lys	Gln	Pro	Asp	Ile	Lys	Asn	Pro	Lys	Leu	Thr	Lys	Asp	His				
385					390					395					400				
Gln	Thr	Thr	Gly	Ser	Pro	Ser	Arg	Lys	Thr	Ile	Thr	Ile	Thr	Val	Lys				
			405					410						415					
Ser	Val	Thr	Ser	Asp	Thr	Ile	His	Ile	Ser	Trp	Lys	Leu	Ala	Leu	Pro				
			420				425						430						
Met	Thr	Ala	Leu	Arg	Leu	Ser	Trp	Leu	Lys	Leu	Gly	His	Ser	Pro	Ala				
	435					440						445							
Phe	Gly	Ser	Ile	Thr	Glu	Thr	Ile	Val	Thr	Gly	Glu	Arg	Ser	Glu	Tyr				
	450				455					460									
Leu	Val	Thr	Ala	Leu	Glu	Pro	Asp	Ser	Pro	Tyr	Lys	Val	Cys	Met	Val				
465				470						475					480				
Pro	Met	Glu	Thr	Ser	Asn	Leu	Tyr	Leu	Phe	Asp	Glu	Thr	Pro	Val	Cys				
			485					490						495					
Ile	Glu	Thr	Glu	Thr	Ala	Pro	Leu	Arg	Met	Tyr	Asn	Pro	Thr	Thr	Thr				
		500					505						510						
Leu	Asn	Arg	Glu	Gln	Glu	Lys	Glu	Pro	Tyr	Lys	Asn	Pro	Asn	Leu	Pro				
	515					520						525							
Leu	Ala	Ala	Ile	Ile	Gly	Gly	Ala	Val	Ala	Leu	Val	Thr	Ile	Ala	Leu				
	530				535						540								
Leu	Ala	Leu	Val	Cys	Trp	Tyr	Val	His	Arg	Asn	Gly	Ser	Leu	Phe	Ser				
545				550						555					560				
Arg	Asn	Cys	Ala	Tyr	Ser	Lys	Gly	Arg	Arg	Arg	Lys	Asp	Asp	Tyr	Ala				

				565						570						575
Glu	Ala	Gly	Thr	Lys	Lys	Asp	Asn	Ser	Ile	Leu	Glu	Ile	Arg	Glu	Thr	
			580					585					590			
Ser	Phe	Gln	Met	Leu	Pro	Ile	Ser	Asn	Glu	Pro	Ile	Ser	Lys	Glu	Glu	
		595					600					605				
Phe	Val	Ile	His	Thr	Ile	Phe	Pro	Pro	Asn	Gly	Met	Asn	Leu	Tyr	Lys	
	610					615					620					
Asn	Asn	His	Ser	Glu	Ser	Ser	Ser	Asn	Arg	Ser	Tyr	Arg	Asp	Ser	Gly	
625					630					635					640	
Ile	Pro	Asp	Ser	Asp	His	Ser	His	Ser								
				645												

<210> 58  
 <211> 660  
 <212> PRT  
 <213> Homo sapiens

<400> 58																
Met	Gly	Leu	Gln	Thr	Thr	Lys	Trp	Pro	Ser	His	Gly	Ala	Phe	Phe	Leu	
1				5				10						15		
Lys	Ser	Trp	Leu	Ile	Ile	Ser	Leu	Gly	Leu	Tyr	Ser	Gln	Val	Ser	Lys	
			20					25					30			
Leu	Leu	Ala	Cys	Pro	Ser	Val	Cys	Arg	Cys	Asp	Arg	Asn	Phe	Val	Tyr	
		35					40					45				
Cys	Asn	Glu	Arg	Ser	Leu	Thr	Ser	Val	Pro	Leu	Gly	Ile	Pro	Glu	Gly	
	50					55					60					
Val	Thr	Val	Leu	Tyr	Leu	His	Asn	Asn	Gln	Ile	Asn	Asn	Ala	Gly	Phe	
	65				70					75					80	
Pro	Ala	Glu	Leu	His	Asn	Val	Gln	Ser	Val	His	Thr	Val	Tyr	Leu	Tyr	
				85				90						95		
Gly	Asn	Gln	Leu	Asp	Glu	Phe	Pro	Met	Asn	Leu	Pro	Lys	Asn	Val	Arg	
			100					105					110			
Val	Leu	His	Leu	Gln	Glu	Asn	Asn	Ile	Gln	Thr	Ile	Ser	Arg	Ala	Ala	
		115					120					125				
Leu	Ala	Gln	Leu	Leu	Lys	Leu	Glu	Glu	Leu	His	Leu	Asp	Asp	Asn	Ser	
	130					135					140					
Ile	Ser	Thr	Val	Gly	Val	Glu	Asp	Gly	Ala	Phe	Arg	Glu	Ala	Ile	Ser	
145					150					155					160	
Leu	Lys	Leu	Leu	Phe	Leu	Ser	Lys	Asn	His	Leu	Ser	Ser	Val	Pro	Val	
				165					170					175		

Gly	Leu	Pro	Val	Asp	Leu	Gln	Glu	Leu	Arg	Val	Asp	Glu	Asn	Arg	Ile	180	185	190	
Ala	Val	Ile	Ser	Asp	Met	Ala	Phe	Gln	Asn	Leu	Thr	Ser	Leu	Glu	Arg	195	200	205	
Leu	Ile	Val	Asp	Gly	Asn	Leu	Leu	Thr	Asn	Lys	Gly	Ile	Ala	Glu	Gly	210	215	220	
Thr	Phe	Ser	His	Leu	Thr	Lys	Leu	Lys	Glu	Phe	Ser	Ile	Val	Arg	Asn	225	230	235	240
Ser	Leu	Ser	His	Pro	Pro	Pro	Asp	Leu	Pro	Gly	Thr	His	Leu	Ile	Arg	245	250	255	
Leu	Tyr	Leu	Gln	Asp	Asn	Gln	Ile	Asn	His	Ile	Pro	Leu	Thr	Ala	Phe	260	265	270	
Ser	Asn	Leu	Arg	Lys	Leu	Glu	Arg	Leu	Asp	Ile	Ser	Asn	Asn	Gln	Leu	275	280	285	
Arg	Met	Leu	Thr	Gln	Gly	Val	Phe	Asp	Asn	Leu	Ser	Asn	Leu	Lys	Gln	290	295	300	
Leu	Thr	Ala	Arg	Asn	Asn	Pro	Trp	Phe	Cys	Asp	Cys	Ser	Ile	Lys	Trp	305	310	315	320
Val	Thr	Glu	Trp	Leu	Lys	Tyr	Ile	Pro	Ser	Ser	Leu	Asn	Val	Arg	Gly	325	330	335	
Phe	Met	Cys	Gln	Gly	Pro	Glu	Gln	Val	Arg	Gly	Met	Ala	Val	Arg	Glu	340	345	350	
Leu	Asn	Met	Asn	Leu	Leu	Ser	Cys	Pro	Thr	Thr	Thr	Pro	Gly	Leu	Pro	355	360	365	
Leu	Phe	Thr	Pro	Ala	Pro	Ser	Thr	Ala	Ser	Pro	Thr	Thr	Gln	Pro	Pro	370	375	380	
Thr	Leu	Ser	Ile	Pro	Asn	Pro	Ser	Arg	Ser	Tyr	Thr	Pro	Pro	Thr	Pro	385	390	395	400
Thr	Thr	Ser	Lys	Leu	Pro	Thr	Ile	Pro	Asp	Trp	Asp	Gly	Arg	Glu	Arg	405	410	415	
Val	Thr	Pro	Pro	Ile	Ser	Glu	Arg	Ile	Gln	Leu	Ser	Ile	His	Phe	Val	420	425	430	
Asn	Asp	Thr	Ser	Ile	Gln	Val	Ser	Trp	Leu	Ser	Leu	Phe	Thr	Val	Met	435	440	445	
Ala	Tyr	Lys	Leu	Thr	Trp	Val	Lys	Met	Gly	His	Ser	Leu	Val	Gly	Gly	450	455	460	
Ile	Val	Gln	Glu	Arg	Ile	Val	Ser	Gly	Glu	Lys	Gln	His	Leu	Ser	Leu	465	470	475	480

Val Asn Leu Glu Pro Arg Ser Thr Tyr Arg Ile Cys Leu Val Pro Leu  
 485 490 495  
 Asp Ala Phe Asn Tyr Arg Ala Val Glu Asp Thr Ile Cys Ser Glu Ala  
 500 505 510  
 Thr Thr His Ala Ser Tyr Leu Asn Asn Gly Ser Asn Thr Ala Ser Ser  
 515 520 525  
 His Glu Gln Thr Thr Ser His Ser Met Gly Ser Pro Phe Leu Leu Ala  
 530 535 540  
 Gly Leu Ile Gly Gly Ala Val Ile Phe Val Leu Val Val Leu Leu Ser  
 545 550 555 560  
 Val Phe Cys Trp His Met His Lys Lys Gly Arg Tyr Thr Ser Gln Lys  
 565 570 575  
 Trp Lys Tyr Asn Arg Gly Arg Arg Lys Asp Asp Tyr Cys Glu Ala Gly  
 580 585 590  
 Thr Lys Lys Asp Asn Ser Ile Leu Glu Met Thr Glu Thr Ser Phe Gln  
 595 600 605  
 Ile Val Ser Leu Asn Asn Asp Gln Leu Leu Lys Gly Asp Phe Arg Leu  
 610 615 620  
 Gln Pro Ile Tyr Thr Pro Asn Gly Gly Ile Asn Tyr Thr Asp Cys His  
 625 630 635 640  
 Ile Pro Asn Asn Met Arg Tyr Cys Asn Ser Ser Val Pro Asp Leu Glu  
 645 650 655  
 His Cys His Thr  
 660  
 <210> 59  
 <211> 674  
 <212> PRT  
 <213> Homo sapiens  
 <400> 59  
 Met Val Val Ala His Pro Thr Ala Thr Ala Thr Thr Thr Pro Thr Ala  
 1 5 10 15  
 Thr Val Thr Ala Thr Val Val Met Thr Thr Ala Thr Met Asp Leu Arg  
 20 25 30  
 Asp Trp Leu Phe Leu Cys Tyr Gly Leu Ile Ala Phe Leu Thr Glu Val  
 35 40 45  
 Ile Asp Ser Thr Thr Cys Pro Ser Val Cys Arg Cys Asp Asn Gly Phe  
 50 55 60  
 Ile Tyr Cys Asn Asp Arg Gly Leu Thr Ser Ile Pro Ala Asp Ile Pro  
 65 70 75 80

Asp	Asp	Ala	Thr	Thr	Leu	Tyr	Leu	Gln	Asn	Asn	Gln	Ile	Asn	Asn	Ala	
				85					90					95		
Gly	Ile	Pro	Gln	Asp	Leu	Lys	Thr	Lys	Val	Asn	Val	Gln	Val	Ile	Tyr	
			100					105					110			
Leu	Tyr	Glu	Asn	Asp	Leu	Asp	Glu	Phe	Pro	Ile	Asn	Leu	Pro	Arg	Ser	
		115					120					125				
Leu	Arg	Glu	Leu	His	Leu	Gln	Asp	Asn	Asn	Val	Arg	Thr	Ile	Ala	Arg	
	130					135					140					
Asp	Ser	Leu	Ala	Arg	Ile	Pro	Leu	Leu	Glu	Lys	Leu	His	Leu	Asp	Asp	
145					150					155					160	
Asn	Ser	Val	Ser	Thr	Val	Ser	Ile	Glu	Glu	Asp	Ala	Phe	Ala	Asp	Ser	
				165					170					175		
Lys	Gln	Leu	Lys	Leu	Leu	Phe	Leu	Ser	Arg	Asn	His	Leu	Ser	Ser	Ile	
			180					185					190			
Pro	Ser	Gly	Leu	Pro	His	Thr	Leu	Glu	Glu	Leu	Arg	Leu	Asp	Asp	Asn	
		195					200					205				
Arg	Ile	Ser	Thr	Ile	Pro	Leu	His	Ala	Phe	Lys	Gly	Leu	Asn	Ser	Leu	
	210					215					220					
Arg	Arg	Leu	Val	Leu	Asp	Gly	Asn	Leu	Leu	Ala	Asn	Gln	Arg	Ile	Ala	
225					230					235					240	
Asp	Asp	Thr	Phe	Ser	Arg	Leu	Gln	Asn	Leu	Thr	Glu	Leu	Ser	Leu	Val	
				245					250					255		
Arg	Asn	Ser	Leu	Ala	Ala	Pro	Pro	Leu	Asn	Leu	Pro	Ser	Ala	His	Leu	
			260					265					270			
Gln	Lys	Leu	Tyr	Leu	Gln	Asp	Asn	Ala	Ile	Ser	His	Ile	Pro	Tyr	Asn	
		275					280					285				
Thr	Leu	Ala	Lys	Met	Arg	Glu	Leu	Glu	Arg	Leu	Asp	Leu	Ser	Asn	Asn	
	290					295					300					
Asn	Leu	Thr	Thr	Leu	Pro	Arg	Gly	Leu	Phe	Asp	Asp	Leu	Gly	Asn	Leu	
305					310					315					320	
Ala	Gln	Leu	Leu	Leu	Arg	Asn	Asn	Pro	Trp	Phe	Cys	Gly	Cys	Asn	Leu	
				325					330					335		
Met	Trp	Leu	Arg	Asp	Trp	Val	Lys	Ala	Arg	Ala	Ala	Val	Val	Asn	Val	
			340					345					350			
Arg	Gly	Leu	Met	Cys	Gln	Gly	Pro	Glu	Lys	Val	Arg	Gly	Met	Ala	Ile	
		355					360					365				
Lys	Asp	Ile	Thr	Ser	Glu	Met	Asp	Glu	Cys	Phe	Glu	Thr	Gly	Pro	Gln	
	370					375					380					

Gly Gly Val Ala Asn Ala Ala Ala Lys Thr Thr Ala Ser Asn His Ala  
 385 390 395 400  
 Ser Ala Thr Thr Pro Gln Gly Ser Leu Phe Thr Leu Lys Ala Lys Arg  
 405 410 415  
 Pro Gly Leu Arg Leu Pro Asp Ser Asn Ile Asp Tyr Pro Met Ala Thr  
 420 425 430  
 Gly Asp Gly Ala Lys Thr Leu Ala Ile His Val Lys Ala Leu Thr Ala  
 435 440 445  
 Asp Ser Ile Arg Ile Thr Trp Lys Ala Thr Leu Pro Ala Ser Ser Phe  
 450 455 460  
 Arg Leu Ser Trp Leu Arg Leu Gly His Ser Pro Ala Val Gly Ser Ile  
 465 470 475 480  
 Thr Glu Thr Leu Val Gln Gly Asp Lys Thr Glu Tyr Leu Leu Thr Ala  
 485 490 495  
 Leu Glu Pro Lys Ser Thr Tyr Ile Ile Cys Met Val Thr Met Glu Thr  
 500 505 510  
 Ser Asn Ala Tyr Val Ala Asp Glu Thr Pro Val Cys Ala Lys Ala Glu  
 515 520 525  
 Thr Ala Asp Ser Tyr Gly Pro Thr Thr Thr Leu Asn Gln Glu Gln Asn  
 530 535 540  
 Ala Gly Pro Met Ala Ser Leu Pro Leu Ala Gly Ile Ile Gly Gly Ala  
 545 550 555 560  
 Val Ala Leu Val Phe Leu Phe Leu Val Leu Gly Ala Ile Cys Trp Tyr  
 565 570 575  
 Val His Gln Ala Gly Glu Leu Leu Thr Arg Glu Arg Ala Tyr Asn Arg  
 580 585 590  
 Gly Ser Arg Glu Lys Asp Asp Tyr Met Glu Ser Gly Thr Lys Lys Asp  
 595 600 605  
 Asn Ser Ile Leu Glu Ile Arg Gly Pro Gly Leu Gln Met Leu Pro Ile  
 610 615 620  
 Asn Pro Tyr Arg Ala Lys Glu Glu Tyr Val Val His Thr Ile Phe Pro  
 625 630 635 640  
 Ser Asn Gly Ser Ser Leu Cys Lys Ala Thr His Thr Ile Gly Tyr Gly  
 645 650 655  
 Thr Thr Arg Gly Tyr Arg Asp Gly Gly Ile Pro Asp Ile Asp Tyr Ser  
 660 665 670  
 Tyr Thr

<210> 60  
 <211> 674  
 <212> PRT  
 <213> Homo sapiens

<400> 60  
 Met Val Val Ala His Pro Thr Ala Thr Ala Thr Thr Thr Pro Thr Ala  
   1                  5                  10                  15  
 Thr Val Thr Ala Thr Val Val Met Thr Thr Ala Thr Met Asp Leu Arg  
                   20                  25                  30  
 Asp Trp Leu Phe Leu Cys Tyr Gly Leu Ile Ala Phe Leu Thr Glu Val  
           35                  40                  45  
 Ile Asp Ser Thr Thr Cys Pro Ser Val Cys Arg Cys Asp Asn Gly Phe  
       50                  55                  60  
 Ile Tyr Cys Asn Asp Arg Gly Leu Thr Ser Ile Pro Ala Asp Ile Pro  
   65                  70                  75                  80  
 Asp Asp Ala Thr Thr Leu Tyr Leu Gln Asn Asn Gln Ile Asn Asn Ala  
                   85                  90                  95  
 Gly Ile Pro Gln Asp Leu Lys Thr Lys Val Asn Val Gln Val Ile Tyr  
           100                  105                  110  
 Leu Tyr Glu Asn Asp Leu Asp Glu Phe Pro Ile Asn Leu Pro Arg Ser  
       115                  120                  125  
 Leu Arg Glu Leu His Leu Gln Asp Asn Asn Val Arg Thr Ile Ala Arg  
   130                  135                  140  
 Asp Ser Leu Ala Arg Ile Pro Leu Leu Glu Lys Leu His Leu Asp Asp  
 145                  150                  155                  160  
 Asn Ser Val Ser Thr Val Ser Ile Glu Glu Asp Ala Phe Ala Asp Ser  
           165                  170                  175  
 Lys Gln Leu Lys Leu Leu Phe Leu Ser Arg Asn His Leu Ser Ser Ile  
       180                  185                  190  
 Pro Ser Gly Leu Pro His Thr Leu Glu Glu Leu Arg Leu Asp Asp Asn  
       195                  200                  205  
 Arg Ile Ser Thr Ile Pro Leu His Ala Phe Lys Gly Leu Asn Ser Leu  
   210                  215                  220  
 Arg Arg Leu Val Leu Asp Gly Asn Leu Leu Ala Asn Gln Arg Ile Ala  
 225                  230                  235                  240  
 Asp Asp Thr Phe Ser Arg Leu Gln Asn Leu Thr Glu Leu Ser Leu Val  
           245                  250                  255  
 Arg Asn Ser Leu Ala Ala Pro Pro Leu Asn Leu Pro Ser Ala His Leu



260					265					270					
Gln	Lys	Leu	Tyr	Leu	Gln	Asp	Asn	Ala	Ile	Ser	His	Ile	Pro	Tyr	Asn
	275						280					285			
Thr	Leu	Ala	Lys	Met	Arg	Glu	Leu	Glu	Arg	Leu	Asp	Leu	Ser	Asn	Asn
	290					295					300				
Asn	Leu	Thr	Thr	Leu	Pro	Arg	Gly	Leu	Phe	Asp	Asp	Leu	Gly	Asn	Leu
305					310					315					320
Ala	Gln	Leu	Leu	Leu	Arg	Asn	Asn	Pro	Trp	Phe	Cys	Gly	Cys	Asn	Leu
					325				330					335	
Met	Trp	Leu	Arg	Asp	Trp	Val	Lys	Ala	Arg	Ala	Ala	Val	Val	Asn	Val
			340					345					350		
Arg	Gly	Leu	Met	Cys	Gln	Gly	Pro	Glu	Lys	Val	Arg	Gly	Met	Ala	Ile
		355					360					365			
Lys	Asp	Ile	Thr	Ser	Glu	Met	Asp	Glu	Cys	Phe	Glu	Thr	Gly	Pro	Gln
	370					375					380				
Gly	Gly	Val	Ala	Asn	Ala	Ala	Ala	Lys	Thr	Thr	Ala	Ser	Asn	His	Ala
385					390					395					400
Ser	Ala	Thr	Thr	Pro	Gln	Gly	Ser	Leu	Phe	Thr	Leu	Lys	Ala	Lys	Arg
				405					410					415	
Pro	Gly	Leu	Arg	Leu	Pro	Asp	Ser	Asn	Ile	Asp	Tyr	Pro	Met	Ala	Thr
			420					425					430		
Gly	Asp	Gly	Ala	Lys	Thr	Leu	Ala	Ile	His	Val	Lys	Ala	Leu	Thr	Ala
		435					440					445			
Asp	Ser	Ile	Arg	Ile	Thr	Trp	Lys	Ala	Thr	Leu	Pro	Ala	Ser	Ser	Phe
	450					455					460				
Arg	Leu	Ser	Trp	Leu	Arg	Leu	Gly	His	Ser	Pro	Ala	Val	Gly	Ser	Ile
465					470					475					480
Thr	Glu	Thr	Leu	Val	Gln	Gly	Asp	Lys	Thr	Glu	Tyr	Leu	Leu	Thr	Ala
			485						490					495	
Leu	Glu	Pro	Lys	Ser	Thr	Tyr	Ile	Ile	Cys	Met	Val	Thr	Met	Glu	Thr
			500					505					510		
Ser	Asn	Ala	Tyr	Val	Ala	Asp	Glu	Thr	Pro	Val	Cys	Ala	Lys	Ala	Glu
		515					520					525			
Thr	Ala	Asp	Ser	Tyr	Gly	Pro	Thr	Thr	Thr	Leu	Asn	Gln	Glu	Gln	Asn
	530					535					540				
Ala	Gly	Pro	Met	Ala	Ser	Leu	Pro	Leu	Ala	Gly	Ile	Ile	Gly	Gly	Ala
545					550					555					560
Val	Ala	Leu	Val	Phe	Leu	Phe	Leu	Val	Leu	Gly	Ala	Ile	Cys	Trp	Tyr

565                      570                      575  
 Val His Gln Ala Gly Glu Leu Leu Thr Arg Glu Arg Ala Tyr Asn Arg  
                          580                      585                      590  
 Gly Ser Arg Glu Lys Asp Asp Tyr Met Glu Ser Gly Thr Lys Lys Asp  
                          595                      600                      605  
 Asn Ser Ile Leu Glu Ile Arg Gly Pro Gly Leu Gln Met Leu Pro Ile  
                          610                      615                      620  
 Asn Pro Tyr Arg Ala Lys Glu Glu Tyr Val Val His Thr Ile Phe Pro  
                          625                      630                      635                      640  
 Ser Asn Gly Ser Ser Leu Cys Lys Ala Thr His Thr Ile Gly Tyr Gly  
                          645                      650                      655  
 Thr Thr Arg Gly Tyr Arg Asp Gly Gly Ile Pro Asp Ile Asp Tyr Ser  
                          660                      665                      670  
 Tyr Thr

<210> 61  
 <211> 246  
 <212> PRT  
 <213> Homo sapiens

<400> 61  
 Pro Met Ala Thr Gly Asp Gly Ala Lys Thr Leu Ala Ile His Val Lys  
   1                          5                          10                          15  
 Ala Leu Thr Ala Asp Ser Ile Arg Ile Thr Trp Lys Ala Thr Leu Pro  
                           20                          25                          30  
 Ala Ser Ser Phe Arg Leu Ser Trp Leu Arg Leu Gly His Ser Pro Ala  
                           35                          40                          45  
 Val Gly Ser Ile Thr Glu Thr Leu Val Gln Gly Asp Lys Thr Glu Tyr  
                           50                          55                          60  
 Leu Leu Thr Ala Leu Glu Pro Lys Ser Thr Tyr Ile Ile Cys Met Val  
                           65                          70                          75                          80  
 Thr Met Glu Thr Ser Asn Ala Tyr Val Ala Asp Glu Thr Pro Val Cys  
                           85                          90                          95  
 Ala Lys Ala Glu Thr Ala Asp Ser Tyr Gly Pro Thr Thr Thr Leu Asn  
                           100                          105                          110  
 Gln Glu Gln Asn Ala Gly Pro Met Ala Ser Leu Pro Leu Ala Gly Ile  
                           115                          120                          125  
 Ile Gly Gly Ala Val Ala Leu Val Phe Leu Phe Leu Val Leu Gly Ala  
                           130                          135                          140

Ile Cys Trp Tyr Val His Gln Ala Gly Glu Leu Leu Thr Arg Glu Arg  
 145 150 155 160  
 Ala Tyr Asn Arg Gly Ser Arg Lys Lys Asp Asp Tyr Met Glu Ser Gly  
 165 170 175  
 Thr Lys Lys Asp Asn Ser Ile Leu Glu Ile Arg Gly Pro Gly Leu Gln  
 180 185 190  
 Met Leu Pro Ile Asn Pro Tyr Arg Ala Lys Glu Glu Tyr Val Val His  
 195 200 205  
 Thr Ile Phe Pro Ser Asn Gly Ser Ser Leu Cys Lys Ala Thr His Thr  
 210 215 220  
 Ile Gly Tyr Gly Thr Thr Arg Gly Tyr Arg Asp Gly Gly Ile Pro Asp  
 225 230 235 240  
 Ile Asp Tyr Ser Tyr Thr  
 245

<210> 62  
 <211> 378  
 <212> PRT  
 <213> Homo sapiens

<400> 62  
 Gly Cys Gly Cys Gly Cys Gly Gly Cys Gly Ala Ala Gly Thr Gly Ala  
 1 5 10 15  
 Ala Thr Thr Thr Gly Cys Thr Gly Gly Ala Cys Ala Cys Gly Thr Cys  
 20 25 30  
 Gly Ala Cys Cys Ala Thr Cys Cys Ala Cys Gly Gly Gly Gly Ala Cys  
 35 40 45  
 Thr Gly Gly Gly Gly Cys Thr Gly Gly Cys Thr Cys Ala Cys Gly Thr  
 50 55 60  
 Ala Thr Cys Cys Gly Gly Cys Thr Cys Ala Thr Gly Gly Gly Thr Gly  
 65 70 75 80  
 Gly Gly Ala Cys Thr Cys Cys Ala Thr Cys Ala Ala Cys Gly Ala Gly  
 85 90 95  
 Gly Thr Gly Gly Ala Cys Gly Ala Gly Thr Cys Cys Thr Thr Cys Cys  
 100 105 110  
 Ala Gly Cys Cys Cys Ala Thr Cys Cys Ala Cys Ala Cys Gly Thr Ala  
 115 120 125  
 Cys Cys Ala Gly Gly Thr Thr Thr Gly Cys Ala Ala Cys Gly Thr Cys  
 130 135 140  
 Ala Thr Gly Ala Gly Cys Cys Cys Cys Ala Ala Cys Cys Ala Gly Ala  
 145 150 155 160

Ala Cys Ala Ala Cys Thr Gly Gly Cys Thr Gly Cys Gly Cys Ala Cys  
 165 170 175  
 Gly Ala Gly Cys Thr Gly Gly Gly Thr Cys Cys Cys Cys Gly Ala  
 180 185 190  
 Gly Ala Cys Gly Gly Cys Gly Cys Cys Cys Gly Gly Cys Gly Cys Gly  
 195 200 205  
 Thr Cys Thr Ala Thr Gly Cys Thr Gly Ala Gly Ala Thr Cys Ala Ala  
 210 215 220  
 Gly Thr Thr Thr Ala Cys Cys Cys Thr Gly Cys Gly Cys Gly Ala Cys  
 225 230 235 240  
 Thr Gly Cys Ala Ala Cys Ala Gly Cys Ala Thr Gly Cys Cys Thr Gly  
 245 250 255  
 Gly Thr Gly Thr Gly Cys Thr Gly Gly Gly Cys Ala Cys Cys Thr Gly  
 260 265 270  
 Cys Ala Ala Gly Gly Ala Gly Ala Cys Cys Thr Thr Cys Ala Ala Cys  
 275 280 285  
 Cys Thr Cys Thr Ala Cys Thr Ala Cys Cys Thr Gly Gly Ala Gly Thr  
 290 295 300  
 Cys Gly Gly Ala Cys Cys Gly Cys Gly Ala Cys Cys Thr Gly Gly Gly  
 305 310 315 320  
 Gly Gly Cys Cys Ala Gly Cys Ala Cys Ala Cys Ala Ala Gly Ala Ala  
 325 330 335  
 Ala Gly Cys Cys Ala Gly Thr Thr Cys Cys Thr Cys Ala Ala Ala Ala  
 340 345 350  
 Thr Cys Gly Ala Cys Ala Cys Cys Ala Thr Thr Gly Cys Gly Gly Cys  
 355 360 365  
 Cys Gly Ala Cys Gly Ala Gly Ala Gly Cys Thr Thr Cys Ala Cys Ala  
 370 375 380  
 Gly Gly Thr Gly Cys Cys Gly Ala Cys Cys Thr Thr Gly Gly Thr Gly  
 385 390 395 400  
 Thr Gly Cys Gly Gly Cys Gly Thr Cys Thr Cys Ala Ala Gly Cys Thr  
 405 410 415  
 Cys Ala Ala Cys Ala Cys Gly Gly Ala Gly Gly Thr Gly Cys Gly Cys  
 420 425 430  
 Ala Gly Thr Gly Thr Gly Gly Gly Thr Cys Cys Cys Cys Thr Cys Ala  
 435 440 445  
 Gly Cys Ala Ala Gly Cys Gly Cys Gly Gly Cys Thr Thr Cys Thr Ala  
 450 455 460

Cys Cys Thr Gly Gly Cys Cys Thr Thr Cys Cys Ala Gly Gly Ala Cys  
 465 470 475 480  
 Ala Thr Ala Gly Gly Thr Gly Cys Cys Thr Gly Cys Cys Thr Gly Gly  
 485 490 495  
 Cys Cys Ala Thr Cys Cys Thr Cys Thr Cys Thr Cys Thr Cys Cys Gly  
 500 505 510  
 Cys Ala Thr Cys Thr Ala Cys Thr Ala Thr Ala Ala Gly Ala Ala Gly  
 515 520 525  
 Thr Gly Cys Cys Cys Thr Gly Cys Cys Ala Thr Gly Gly Thr Gly Cys  
 530 535 540  
 Gly Cys Ala Ala Thr Cys Thr Gly Gly Cys Thr Gly Cys Cys Thr Thr  
 545 550 555 560  
 Cys Thr Cys Gly Gly Ala Gly Gly Cys Ala Gly Thr Gly Ala Cys Gly  
 565 570 575  
 Gly Gly Gly Gly Cys Cys Gly Ala Cys Thr Cys Gly Thr Cys Cys Thr  
 580 585 590  
 Cys Ala Cys Thr Gly Gly Thr Gly Gly Ala Gly Gly Thr Gly Ala Gly  
 595 600 605  
 Gly Gly Gly Cys Cys Ala Gly Thr Gly Cys Gly Thr Gly Cys Gly Gly  
 610 615 620  
 Cys Ala Cys Thr Cys Ala Gly Ala Gly Gly Ala Gly Cys Gly Gly Gly  
 625 630 635 640  
 Ala Cys Ala Cys Ala Cys Cys Cys Ala Ala Gly Ala Thr Gly Thr Ala  
 645 650 655  
 Cys Thr Gly Cys Ala Gly Cys Gly Cys Gly Gly Ala Gly Gly Gly Cys  
 660 665 670  
 Gly Ala Gly Thr Gly Gly Cys Thr Cys Gly Thr Gly Cys Cys Cys Ala  
 675 680 685  
 Thr Cys Gly Gly Cys Ala Ala Ala Thr Gly Cys Gly Thr Gly Thr Gly  
 690 695 700  
 Cys Ala Gly Thr Gly Cys Cys Gly Gly Cys Thr Ala Cys Gly Ala Gly  
 705 710 715 720  
 Gly Ala Gly Cys Gly Gly Cys Gly Gly Gly Ala Thr Gly Cys Cys Thr  
 725 730 735  
 Gly Thr Gly Thr Gly Gly Cys Cys Thr Gly Thr Gly Ala Gly Cys Thr  
 740 745 750  
 Gly Gly Gly Cys Thr Thr Cys Thr Ala Cys Ala Ala Gly Thr Cys Ala  
 755 760 765

Gly Cys Cys Cys Cys Thr Gly Gly Gly Gly Ala Cys Cys Ala Gly Cys  
 770 775 780  
 Thr Gly Thr Gly Thr Gly Cys Cys Cys Gly Cys Thr Gly Cys Cys Cys  
 785 790 795 800  
 Thr Cys Cys Cys Cys Ala Cys Ala Gly Cys Cys Ala Cys Thr Cys Cys  
 805 810 815  
 Gly Cys Ala Gly Cys Thr Cys Cys Ala Gly Cys Cys Gly Cys Cys Cys  
 820 825 830  
 Ala Ala Gly Cys Cys Thr Gly Cys Cys Ala Cys Thr Gly Thr Gly Ala  
 835 840 845  
 Cys Cys Thr Cys Ala Gly Cys Thr Ala Cys Thr Ala Cys Cys Gly Thr  
 850 855 860  
 Gly Cys Ala Gly Cys Cys Cys Thr Gly Gly Ala Cys Cys Cys Gly Cys  
 865 870 875 880  
 Cys Gly Thr Cys Cys Thr Cys Ala Gly Cys Cys Thr Gly Cys Ala Cys  
 885 890 895  
 Cys Cys Gly Gly Cys Cys Ala Cys Cys Cys Thr Cys Gly Gly Cys Ala  
 900 905 910  
 Cys Cys Ala Gly Thr Gly Ala Ala Cys Cys Thr Gly Ala Thr Cys Thr  
 915 920 925  
 Cys Cys Ala Gly Thr Gly Thr Gly Ala Ala Thr Gly Gly Gly Ala Cys  
 930 935 940  
 Ala Thr Cys Ala Gly Thr Gly Ala Cys Thr Cys Thr Gly Gly Ala Gly  
 945 950 955 960  
 Thr Gly Gly Gly Cys Cys Cys Thr Cys Cys Cys Thr Gly Gly  
 965 970 975  
 Ala Cys Cys Cys Ala Gly Gly Thr Gly Gly Cys Cys Gly Cys Ala Gly  
 980 985 990  
 Thr Gly Ala Cys Ala Thr Cys Ala Cys Cys Thr Ala Cys Ala Ala Thr  
 995 1000 1005  
 Gly Cys Cys Gly Thr Gly Thr Gly Cys Cys Gly Cys  
 1010 1015 1020

<210> 63  
 <211> 338  
 <212> PRT  
 <213> Gallus gallus

<400> 63  
 Ala Arg Gly Glu Val Asn Leu Leu Asp Thr Ser Thr Ile His Gly Asp

1	5	10	15
Trp Gly Trp Leu Thr Tyr Pro Ala His Gly Trp Asp Ser Ile Asn Glu	20	25	30
Val Asp Glu Ser Phe Gln Pro Ile His Thr Tyr Gln Val Cys Asn Val	35	40	45
Met Ser Pro Asn Gln Asn Asn Trp Leu Arg Thr Ser Trp Val Pro Arg	50	55	60
Asp Gly Ala Arg Arg Val Tyr Ala Glu Ile Lys Phe Thr Leu Arg Asp	65	70	80
Cys Asn Ser Met Pro Gly Val Leu Gly Thr Cys Lys Glu Thr Phe Asn	85	90	95
Leu Tyr Tyr Leu Glu Ser Asp Arg Asp Leu Gly Ala Ser Thr Gln Glu	100	105	110
Ser Gln Phe Leu Lys Ile Asp Thr Ile Ala Ala Asp Glu Ser Phe Thr	115	120	125
Gly Ala Asp Leu Gly Val Arg Arg Leu Lys Leu Asn Thr Glu Val Arg	130	135	140
Ser Val Gly Pro Leu Ser Lys Arg Gly Phe Tyr Leu Ala Phe Gln Asp	145	150	160
Ile Gly Ala Cys Leu Ala Ile Leu Ser Leu Arg Ile Tyr Tyr Lys Lys	165	170	175
Cys Pro Ala Met Val Arg Asn Leu Ala Ala Phe Ser Glu Ala Val Thr	180	185	190
Gly Ala Asp Ser Ser Ser Leu Val Glu Val Arg Gly Gln Cys Val Arg	195	200	205
His Ser Glu Glu Arg Asp Thr Pro Lys Met Tyr Cys Ser Ala Glu Gly	210	215	220
Glu Trp Leu Val Pro Ile Gly Lys Cys Val Cys Ser Ala Gly Tyr Glu	225	230	240
Glu Arg Arg Asp Ala Cys Val Ala Cys Glu Leu Gly Phe Tyr Lys Ser	245	250	255
Ala Pro Gly Asp Gln Leu Cys Ala Arg Cys Pro Pro His Ser His Ser	260	265	270
Ala Ala Pro Ala Ala Gln Ala Cys His Cys Asp Leu Ser Tyr Tyr Arg	275	280	285
Ala Ala Leu Asp Pro Pro Ser Ser Ala Cys Thr Arg Pro Pro Ser Ala	290	295	300
Pro Val Asn Leu Ile Ser Ser Val Asn Gly Thr Ser Val Thr Leu Glu			





Ala Cys Thr Gly Thr Cys Ala Cys Gly Gly Cys Cys Ala Cys Cys Gly  
 50 55 60  
 Thr Thr Gly Thr Gly Ala Thr Gly Ala Cys Cys Ala Cys Gly Gly Cys  
 65 70 75 80  
 Cys Ala Cys Cys Ala Thr Gly Gly Ala Cys Cys Thr Gly Cys Gly Gly  
 85 90 95  
 Gly Ala Cys Thr Gly Gly Cys Thr Gly Thr Thr Cys Cys Thr Cys Thr  
 100 105 110  
 Gly Cys Thr Ala Cys Gly Gly Gly Cys Thr Cys Ala Thr Cys Gly Cys  
 115 120 125  
 Cys Thr Thr Cys Cys Thr Gly Ala Cys Gly Gly Ala Gly Gly Thr Cys  
 130 135 140  
 Ala Thr Cys Gly Ala Cys Ala Gly Cys Ala Cys Cys Ala Cys Cys Thr  
 145 150 155 160  
 Gly Cys Cys Cys Cys Thr Cys Gly Gly Thr Gly Thr Gly Cys Cys Gly  
 165 170 175  
 Cys Thr Gly Cys Gly Ala Cys Ala Ala Cys Gly Gly Cys Thr Thr Cys  
 180 185 190  
 Ala Thr Cys Thr Ala Cys Thr Gly Cys Ala Ala Cys Gly Ala Cys Cys  
 195 200 205  
 Gly Gly Gly Gly Ala Cys Thr Cys Ala Cys Ala Thr Cys Cys Ala Thr  
 210 215 220  
 Cys Cys Cys Cys Gly Cys Ala Gly Ala Thr Ala Thr Cys Cys Cys Thr  
 225 230 235 240  
 Gly Ala Thr Gly Ala Thr Gly Cys Cys Ala Cys Cys Ala Cys Cys Cys  
 245 250 255  
 Thr Cys Thr Ala Cys Cys Thr Gly Cys Ala Gly Ala Ala Cys Ala Ala  
 260 265 270  
 Cys Cys Ala Gly Ala Thr Cys Ala Ala Cys Ala Ala Cys Gly Cys Cys  
 275 280 285  
 Gly Gly Cys Ala Thr Cys Cys Cys Cys Cys Ala Gly Gly Ala Cys Cys  
 290 295 300  
 Thr Cys Ala Ala Gly Ala Cys Cys Ala Ala Gly Gly Thr Cys Ala Ala  
 305 310 315 320  
 Cys Gly Thr Gly Cys Ala Gly Gly Thr Cys Ala Thr Cys Thr Ala Cys  
 325 330 335  
 Cys Thr Ala Thr Ala Cys Gly Ala Gly Ala Ala Thr Gly Ala Cys Cys  
 340 345 350

Thr Gly Gly Ala Thr Gly Ala Gly Thr Thr Cys Cys Cys Cys Ala Thr  
 355 360 365  
 Cys Ala Ala Cys Cys Thr Gly Cys Cys Cys Cys Gly Cys Thr Cys Cys  
 370 375 380  
 Cys Thr Cys Cys Gly Gly Gly Ala Gly Cys Thr Gly Cys Ala Cys Cys  
 385 390 395 400  
 Thr Gly Cys Ala Gly Gly Ala Cys Ala Ala Cys Ala Ala Thr Gly Thr  
 405 410 415  
 Gly Cys Gly Cys Ala Cys Cys Ala Thr Thr Gly Cys Cys Ala Gly Gly  
 420 425 430  
 Gly Ala Cys Thr Cys Gly Cys Thr Gly Gly Cys Cys Cys Gly Cys Ala  
 435 440 445  
 Thr Cys Cys Cys Gly Cys Thr Gly Cys Thr Gly Gly Ala Gly Ala Ala  
 450 455 460  
 Gly Cys Thr Gly Cys Ala Cys Cys Thr Gly Gly Ala Thr Gly Ala Cys  
 465 470 475 480  
 Ala Ala Cys Thr Cys Cys Gly Thr Gly Thr Cys Cys Ala Cys Cys Gly  
 485 490 495  
 Thr Cys Ala Gly Cys Ala Thr Thr Gly Ala Gly Gly Ala Gly Gly Ala  
 500 505 510  
 Cys Gly Cys Cys Thr Thr Cys Gly Cys Cys Gly Ala Cys Ala Gly Cys  
 515 520 525  
 Ala Ala Ala Cys Ala Gly Cys Thr Cys Ala Ala Gly Cys Thr Gly Cys  
 530 535 540  
 Thr Cys Thr Thr Cys Cys Thr Gly Ala Gly Cys Cys Gly Gly Ala Ala  
 545 550 555 560  
 Cys Cys Ala Cys Cys Thr Gly Ala Gly Cys Ala Gly Cys Ala Thr Cys  
 565 570 575  
 Cys Cys Cys Thr Cys Gly Gly Gly Gly Cys Thr Gly Cys Cys Gly Cys  
 580 585 590  
 Ala Cys Ala Cys Gly Cys Thr Gly Gly Ala Gly Gly Ala Gly Cys Thr  
 595 600 605  
 Gly Cys Gly Gly Cys Thr Gly Gly Ala Thr Gly Ala Cys Ala Ala Cys  
 610 615 620  
 Cys Gly Cys Ala Thr Cys Thr Cys Cys Ala Cys Cys Ala Thr Cys Cys  
 625 630 635 640  
 Cys Gly Cys Thr Gly Cys Ala Thr Gly Cys Cys Thr Thr Cys Ala Ala  
 645 650 655

Gly Gly Gly Cys Cys Thr Cys Ala Ala Cys Ala Gly Cys Cys Thr Gly  
 660 665 670  
 Cys Gly Gly Cys Gly Cys Cys Thr Gly Gly Thr Gly Cys Thr Gly Gly  
 675 680 685  
 Ala Cys Gly Gly Thr Ala Ala Cys Cys Thr Gly Cys Thr Gly Gly Cys  
 690 695 700  
 Cys Ala Ala Cys Cys Ala Gly Cys Gly Cys Ala Thr Cys Gly Cys Cys  
 705 710 715 720  
 Gly Ala Cys Gly Ala Cys Ala Cys Cys Thr Thr Cys Ala Gly Cys Cys  
 725 730 735  
 Gly Cys Cys Thr Ala Cys Ala Gly Ala Ala Cys Cys Thr Cys Ala Cys  
 740 745 750  
 Ala Gly Ala Gly Cys Thr Cys Thr Cys Gly Cys Thr Gly Gly Thr Gly  
 755 760 765  
 Cys Gly Cys Ala Ala Thr Thr Cys Gly Cys Thr Gly Gly Cys Cys Gly  
 770 775 780  
 Cys Gly Cys Cys Ala Cys Cys Cys Cys Thr Cys Thr Ala Cys Cys Thr  
 785 790 795 800  
 Gly Cys Ala Gly Gly Ala Cys Ala Ala Thr Gly Cys Cys Ala Thr Cys  
 805 810 815  
 Ala Gly Cys Cys Ala Cys Ala Thr Cys Cys Cys Cys Thr Ala Cys Ala  
 820 825 830  
 Ala Cys Ala Cys Gly Cys Thr Gly Gly Cys Cys Ala Ala Gly Ala Thr  
 835 840 845  
 Gly Cys Gly Thr Gly Ala Gly Cys Thr Gly Gly Ala Gly Cys Gly Gly  
 850 855 860  
 Cys Thr Gly Gly Ala Cys Cys Thr Gly Thr Cys Cys Ala Ala Cys Ala  
 865 870 875 880  
 Ala Cys Ala Ala Cys Cys Thr Gly Ala Cys Cys Ala Cys Gly Cys Thr  
 885 890 895  
 Gly Cys Cys Cys Cys Gly Cys Gly Gly Cys Cys Thr Gly Thr Thr Cys  
 900 905 910  
 Gly Ala Cys Gly Ala Cys Cys Thr Gly Gly Gly Gly Ala Ala Cys Cys  
 915 920 925  
 Thr Gly Gly Cys Cys Cys Ala Gly Cys Thr Gly Cys Thr Gly Cys Thr  
 930 935 940  
 Cys Ala Gly Gly Ala Ala Cys Ala Ala Cys Cys Cys Thr Thr Gly Gly  
 945 950 955 960

Thr Thr Thr Thr Gly Thr Gly Gly Cys Thr Gly Cys Ala Ala Cys Cys  
 965 970 975  
 Thr Cys Ala Thr Gly Thr Gly Gly Cys Thr Gly Cys Gly Gly Gly Ala  
 980 985 990  
 Cys Thr Gly Gly Gly Thr Gly Ala Ala Gly Gly Cys Ala Cys Gly Gly  
 995 1000 1005  
 Gly Cys Gly Gly Cys Cys Gly Thr Gly Gly Thr Cys Ala Ala Cys Gly  
 1010 1015 1020  
 Thr Gly Cys Gly Gly Gly Gly Cys Cys Thr Cys Ala Thr Gly Thr Gly  
 1025 1030 1035 1040  
 Cys Cys Ala Gly Gly Gly Cys Cys Cys Thr Gly Ala Gly Ala Ala Gly  
 1045 1050 1055  
 Gly Thr Cys Cys Gly Gly Gly Gly Cys Ala Thr Gly Gly Cys Cys Ala  
 1060 1065 1070  
 Thr Cys Ala Ala Gly Gly Ala Cys Ala Thr Thr Ala Cys Cys Ala Gly  
 1075 1080 1085  
 Cys Gly Ala Gly Gly Thr Gly Gly Ala Gly Ala Gly Thr Gly Thr Thr  
 1090 1095 1100  
 Thr Thr Gly Ala Gly Ala Cys Gly Gly Gly Cys Gly Cys Cys Gly Cys  
 1105 1110 1115 1120  
 Ala Gly Gly Gly Cys Gly Gly Cys Gly Thr Gly Gly Cys Cys Ala Ala  
 1125 1130 1135  
 Thr Gly Cys Gly Gly Cys Thr Gly Cys Cys Ala Ala Gly Ala Cys Cys  
 1140 1145 1150  
 Ala Cys Gly Gly Cys Cys Ala Gly Cys Ala Ala Cys Cys Ala Cys Gly  
 1155 1160 1165  
 Cys Cys Thr Cys Thr Gly Cys Cys Ala Cys Cys Ala Cys Gly Cys Cys  
 1170 1175 1180  
 Cys Cys Ala Gly Gly Gly Thr Thr Cys Cys Cys Thr Gly Thr Thr Thr  
 1185 1190 1195 1200  
 Ala Cys Cys Cys Thr Cys Ala Ala Gly Gly Cys Cys Ala Ala Ala Ala  
 1205 1210 1215  
 Gly Gly Cys Cys Ala Gly Gly Gly Cys Thr Gly Cys Gly Cys Cys Thr  
 1220 1225 1230  
 Cys Cys Cys Cys Gly Ala Cys Thr Cys Cys Ala Ala Cys Ala Thr Thr  
 1235 1240 1245  
 Gly Ala Cys Thr Ala Cys Cys Cys Cys Ala Thr Gly Gly Cys Cys Ala  
 1250 1255 1260

Cys Gly Gly Gly Thr Gly Ala Thr Gly Gly Cys Gly Cys Cys Ala Ala  
 1265 1270 1275 1280  
 Gly Ala Cys Cys Cys Thr Gly Gly Cys Cys Ala Thr Cys Cys Ala Cys  
 1285 1290 1295  
 Gly Thr Gly Ala Ala Gly Gly Cys Cys Cys Thr Gly Ala Cys Gly Gly  
 1300 1305 1310  
 Cys Ala Gly Ala Cys Thr Cys Cys Ala Thr Cys Cys Gly Cys Ala Thr  
 1315 1320 1325  
 Cys Ala Cys Gly Thr Gly Gly Ala Ala Gly Gly Cys Cys Ala Cys Gly  
 1330 1335 1340  
 Cys Thr Cys Cys Cys Cys Gly Cys Cys Thr Cys Cys Thr Cys Thr Thr  
 1345 1350 1355 1360  
 Thr Cys Cys Gly Gly Cys Thr Cys Ala Gly Thr Thr Gly Gly Cys Thr  
 1365 1370 1375  
 Gly Cys Gly Cys Cys Thr Gly Gly Gly Cys Cys Ala Cys Ala Gly Cys  
 1380 1385 1390  
 Cys Cys Ala Gly Cys Cys Gly Thr Gly Gly Gly Cys Thr Cys Cys Ala  
 1395 1400 1405  
 Thr Cys Ala Cys Gly Gly Ala Gly Ala Cys Cys Thr Thr Gly Gly Thr  
 1410 1415 1420  
 Gly Cys Ala Gly Gly Gly Gly Gly Ala Cys Ala Ala Gly Ala Cys Ala  
 1425 1430 1435 1440  
 Gly Ala Gly Thr Ala Cys Cys Thr Gly Cys Thr Gly Ala Cys Ala Gly  
 1445 1450 1455  
 Cys Cys Cys Thr Gly Gly Ala Gly Cys Cys Cys Ala Ala Gly Thr Cys  
 1460 1465 1470  
 Cys Ala Cys Cys Thr Ala Cys Ala Thr Cys Ala Thr Cys Thr Gly Cys  
 1475 1480 1485  
 Ala Thr Gly Gly Thr Cys Ala Cys Cys Ala Thr Gly Gly Ala Gly Ala  
 1490 1495 1500  
 Cys Cys Ala Gly Cys Ala Ala Thr Gly Cys Cys Thr Ala Cys Gly Thr  
 1505 1510 1515 1520  
 Ala Gly Cys Thr Gly Ala Thr Gly Ala Gly Ala Cys Ala Cys Cys Cys  
 1525 1530 1535  
 Gly Thr Gly Thr Gly Thr Gly Cys Cys Ala Ala Gly Gly Cys Ala Gly  
 1540 1545 1550  
 Ala Gly Ala Cys Ala Gly Cys Cys Gly Ala Cys Ala Gly Cys Thr Ala  
 1555 1560 1565

Thr Gly Gly Cys Cys Cys Thr Ala Cys Cys Ala Cys Cys Ala Cys Ala  
 1570 1575 1580

Cys Thr Cys Ala Ala Cys Cys Ala Gly Gly Ala Gly Cys Ala Gly Ala  
 1585 1590 1595 1600

Ala Cys Gly Cys Thr Gly Gly Cys Cys Cys Cys Ala Thr Gly Gly Cys  
 1605 1610 1615

Gly Ala Gly Cys Cys Thr Gly Cys Cys Cys Cys Thr Gly Gly Cys Gly  
 1620 1625 1630

Gly Gly Cys Ala Thr Cys Ala Thr Cys Gly Gly Cys Gly Gly Gly Gly  
 1635 1640 1645

Cys Ala Gly Thr Gly Gly Cys Thr Cys Thr Gly Gly Thr Cys Thr Thr  
 1650 1655 1660

Cys Cys Thr Cys Thr Thr Cys Cys Thr Gly Gly Thr Cys Cys Thr Gly  
 1665 1670 1675 1680

Gly Gly Gly Gly Cys Cys Ala Thr Cys Thr Gly Cys Thr Gly Gly Thr  
 1685 1690 1695

Ala Cys Gly Thr Gly Cys Ala Cys Cys Ala Gly Gly Cys Thr Gly Gly  
 1700 1705 1710

Cys Gly Ala Gly Cys Thr Gly Cys Thr Gly Ala Cys Cys Cys Gly Gly  
 1715 1720 1725

Gly Ala Gly Ala Gly Gly Gly Cys Cys Thr Ala Cys Ala Ala Cys Cys  
 1730 1735 1740

Gly Gly Gly Gly Cys Ala Gly Cys Ala Gly Gly Ala Ala Ala Ala Ala  
 1745 1750 1755 1760

Gly Gly Ala Thr Gly Ala Cys Thr Ala Thr Ala Thr Gly Gly Ala Gly  
 1765 1770 1775

Thr Cys Ala Gly Gly Gly Ala Cys Cys Ala Ala Gly Ala Ala Gly Gly  
 1780 1785 1790

Ala Thr Ala Ala Cys Thr Cys Cys Ala Thr Cys Cys Thr Gly Gly Ala  
 1795 1800 1805

Ala Ala Thr Cys Cys Gly Cys Gly Gly Cys Cys Cys Thr Gly Gly Gly  
 1810 1815 1820

Cys Thr Gly Cys Ala Gly Ala Thr Gly Cys Thr Gly Cys Cys Cys Ala  
 1825 1830 1835 1840

Thr Cys Ala Ala Cys Cys Cys Gly Thr Ala Cys Cys Gly Cys Gly Cys  
 1845 1850 1855

Cys Ala Ala Ala Gly Ala Ala Gly Ala Gly Thr Ala Cys Gly Thr Gly  
 1860 1865 1870

Gly Thr Cys Cys Ala Cys Ala Cys Thr Ala Thr Cys Thr Thr Cys Cys  
 1875 1880 1885

Cys Cys Thr Cys Cys Ala Ala Cys Gly Gly Cys Ala Gly Cys Ala Gly  
 1890 1895 1900

Cys Cys Thr Cys Thr Gly Cys Ala Ala Gly Gly Cys Cys Ala Cys Ala  
 1905 1910 1915 1920

Cys Ala Cys Ala Cys Cys Ala Thr Thr Gly Gly Cys Thr Ala Cys Gly  
 1925 1930 1935

Gly Cys Ala Cys Cys Ala Cys Gly Cys Gly Gly Gly Cys Thr Ala  
 1940 1945 1950

Cys Cys Gly Gly Gly Ala Cys Gly Gly Cys Gly Gly Cys Ala Thr Cys  
 1955 1960 1965

Cys Cys Cys Gly Ala Cys Ala Thr Ala Gly Ala Cys Thr Ala Cys Thr  
 1970 1975 1980

Cys Cys Thr Ala Cys Cys Ala  
 1985 1990

<210> 65  
 <211> 1020  
 <212> DNA  
 <213> Homo sapiens

<400> 65  
 gcgcgcggcg aagtgaattt gctggacacg tcgaccatcc acgggggactg gggctggctc 60  
 acgtatccgg ctcattgggtg ggactccatc aacgaggtgg acgagtcctt ccagcccac 120  
 cacacgtacc aggtttgcaa cgtcatgagc cccaaccaga acaactggct gcgcacgagc 180  
 tgggtccccc gagacggcgc ccggcgcgct tatgctgaga tcaagtttac cctgcgcgac 240  
 tgcaacagca tgcttggtgt gctgggcacc tgcaaggaga ccttcaacct ctactacctg 300  
 gagtcggacc gcgacctggg ggccagcaca caagaaagcc agttcctcaa aatcgacacc 360  
 attgcggccg acgagagctt cacaggtgcc gaccttgggtg tgcggcgtct caagctcaac 420  
 acggaggtgc gcagtgtggg tcccctcagc aagcgcggct tctacctggc cttccaggac 480  
 ataggtgcct gcctggccat cctctctctc cgcattctact ataagaagtg ccctgccatg 540  
 gtgcgcaatc tggtgcctt ctgggaggca gtgacggggg ccgactcgtc ctcactggtg 600  
 gaggtgaggg gccagtgcgt gcggcactca gaggagcggg acacacccaa gatgtactgc 660  
 agcgcggagg gcgagtggct cgtgcccacg ggcaaagtgc tgtgcagtgc cggctacgag 720  
 gagcggcggg atgcctgtgt ggctgtgag ctgggcttct acaagtcagc ccctggggac 780  
 cagctgtgtg ccgctgccc tcccacagc cactccgcag ctccagccgc ccaagcctgc 840  
 cactgtgacc tcagctacta ccgtgcagcc ctggaccgc cgtcctcagc ctgcaccggg 900  
 ccaccctcgg caccagtga cctgatctcc agtgtgaatg ggacatcagt gactctggag 960  
 tgggcccctc ccctggaccc aggtggccgc agtgacatca cctacaatgc cgtgtgccgc 1020

<210> 66  
 <211> 515  
 <212> PRT  
 <213> Homo sapiens

<400> 66  
 Ala Arg Gly Glu Val Asn Leu Leu Asp Thr Ser Thr Ile His Gly Asp

1		5		10		15									
Trp	Gly	Trp	Leu	Thr	Tyr	Pro	Ala	His	Gly	Trp	Asp	Ser	Ile	Asn	Glu
		20						25					30		
Val	Asp	Glu	Ser	Phe	Gln	Pro	Ile	His	Thr	Tyr	Gln	Val	Cys	Asn	Val
		35					40					45			
Met	Ser	Pro	Asn	Gln	Asn	Asn	Trp	Leu	Arg	Thr	Ser	Trp	Val	Pro	Arg
	50					55					60				
Asp	Gly	Ala	Arg	Arg	Val	Tyr	Ala	Glu	Ile	Lys	Phe	Thr	Leu	Arg	Asp
65					70					75					80
Cys	Asn	Ser	Met	Pro	Gly	Val	Leu	Gly	Thr	Cys	Lys	Glu	Thr	Phe	Asn
				85					90					95	
Leu	Tyr	Tyr	Leu	Glu	Ser	Asp	Arg	Asp	Leu	Gly	Ala	Ser	Thr	Gln	Glu
			100					105						110	
Ser	Gln	Phe	Leu	Lys	Ile	Asp	Thr	Ile	Ala	Ala	Asp	Glu	Ser	Phe	Thr
		115					120					125			
Gly	Ala	Asp	Leu	Gly	Val	Arg	Arg	Leu	Lys	Leu	Asn	Thr	Glu	Val	Arg
	130					135					140				
Ser	Val	Gly	Pro	Leu	Ser	Lys	Arg	Gly	Phe	Tyr	Leu	Ala	Phe	Gln	Asp
145					150					155					160
Ile	Gly	Ala	Cys	Leu	Ala	Ile	Leu	Ser	Leu	Arg	Ile	Tyr	Tyr	Lys	Lys
			165						170					175	
Cys	Pro	Ala	Met	Val	Arg	Asn	Leu	Ala	Ala	Phe	Ser	Glu	Ala	Val	Thr
			180					185					190		
Gly	Ala	Asp	Ser	Ser	Ser	Leu	Val	Glu	Val	Arg	Gly	Gln	Cys	Val	Arg
		195					200					205			
His	Ser	Glu	Glu	Arg	Asp	Thr	Pro	Lys	Met	Tyr	Cys	Ser	Ala	Glu	Gly
	210					215					220				
Glu	Trp	Leu	Val	Pro	Ile	Gly	Lys	Cys	Val	Cys	Ser	Ala	Gly	Tyr	Glu
225					230					235					240
Glu	Arg	Arg	Asp	Ala	Cys	Val	Ala	Cys	Glu	Leu	Gly	Phe	Tyr	Lys	Ser
			245						250					255	
Ala	Pro	Gly	Asp	Gln	Leu	Cys	Ala	Arg	Cys	Pro	Pro	His	Ser	His	Ser
			260					265					270		
Ala	Ala	Pro	Ala	Ala	Gln	Ala	Cys	His	Cys	Asp	Leu	Ser	Tyr	Tyr	Arg
		275					280					285			
Ala	Ala	Leu	Asp	Pro	Pro	Ser	Ser	Ala	Cys	Thr	Arg	Pro	Pro	Ser	Ala
	290					295					300				
Pro	Val	Asn	Leu	Ile	Ser	Ser	Val	Asn	Gly	Thr	Ser	Val	Thr	Leu	Glu





```

ctgctcttcc tgagccggaa ccacctgagc agcatccccct cggggctgcc gcacacgctg 600
gaggagctgc ggctggatga caaccgcatc tccaccatcc cgctgcatgc cttcaagggc 660
ctcaacagcc tgccggcgcc ggtgctggac ggtaacctgc tggccaacca gcgcatcgcc 720
gacgacacct tcagccgcct acagaacctc acagagctct cgctgggtgc caattcgctg 780
gccgcgccac cctctacct gcaggacaat gccatcagcc acatccccta caacacgctg 840
gccaagatgc gtgagctgga gcggctggac ctgtccaaca acaacctgac cacgctgccc 900
cgcgccctgt tcgacgacct ggggaacctg gcccagctgc tgctcaggaa caacccttgg 960
ttttgtggct gcaacctcat gtggctgcgg gactgggtga aggcacgggc ggccgtggtc 1020
aacgtgcggg gcctcatgtg ccagggccct gagaagggtcc ggggcatggc catcaaggac 1080
attaccagcg aggtggagag tgttttgaga cgggcgcgcg agggcgcgct ggccaatgcy 1140
gctgccaaga ccacggccag caaccacgcc tctgccacca cgcccagggt ttccctgttt 1200
accctcaagg ccaaaaggcc agggctgcgc ctccccgact ccaacattga ctaccccatg 1260
gccacgggtg atggcgccaa gaccctggcc atccacgtga aggcctgac ggcagactcc 1320
atccgcatca cgtggaaggc cacgtcccc gcctcctctt tccggctcag ttggctgcgc 1380
ctgggccaca gccagccgt gggctccatc acggagacct tggcgagggt ggacaagaca 1440
gagtacctgc tgacagccct ggagcccaag tccacctaca tcatctgcat ggtcaccatg 1500
gagaccagca atgcctacgt agctgatgag acaccctgtg gtgccaaggc agagacagcc 1560
gacagctatg gccctaccac cacactcaac caggagcaga acgctggccc catggcgagc 1620
ctgcccctgg cgggcatcat cggcggggca gtggctctgg tctcctctt cctggctctg 1680
ggggccatct gctggtacgt gcaccaggct ggcgagctgc tgaccgggga gagggcctac 1740
aaccggggca gcaggaaaaa ggatgactat atggagtcag ggaccaagaa ggataactcc 1800
atcctggaaa tccgcggccc tgggctgcag atgctgccc tcaaccgta ccgcgcaaaa 1860
gaagagtacg tgggtccacac tatcttcccc tccaacggca gcagcctctg caaggccaca 1920
cacaccattg gctacggcac cacgcggggc taccgggacg gcggcatccc cgacatagac 1980
tactcctaca ca

```

```

<210> 68
<211> 664
<212> PRT
<213> Homo sapiens

```

```

<400> 68
Met Val Val Ala His Pro Thr Ala Thr Ala Thr Thr Thr Pro Thr Ala
  1                      5                      10                      15

Thr Val Thr Ala Thr Val Val Met Thr Thr Ala Thr Met Asp Leu Arg
          20                      25                      30

Asp Trp Leu Phe Leu Cys Tyr Gly Leu Ile Ala Phe Leu Thr Glu Val
          35                      40                      45

Ile Asp Ser Thr Thr Cys Pro Ser Val Cys Arg Cys Asp Asn Gly Phe
          50                      55                      60

Ile Tyr Cys Asn Asp Arg Gly Leu Thr Ser Ile Pro Ala Asp Ile Pro
          65                      70                      75                      80

Asp Asp Ala Thr Thr Leu Tyr Leu Gln Asn Asn Gln Ile Asn Asn Ala
          85                      90                      95

Gly Ile Pro Gln Asp Leu Lys Thr Lys Val Asn Val Gln Val Ile Tyr
          100                      105                      110

Leu Tyr Glu Asn Asp Leu Asp Glu Phe Pro Ile Asn Leu Pro Arg Ser
          115                      120                      125

```

Leu Arg Glu Leu His Leu Gln Asp Asn Asn Val Arg Thr Ile Ala Arg  
 130 135 140  
 Asp Ser Leu Ala Arg Ile Pro Leu Leu Glu Lys Leu His Leu Asp Asp  
 145 150 155 160  
 Asn Ser Val Ser Thr Val Ser Ile Glu Glu Asp Ala Phe Ala Asp Ser  
 165 170 175  
 Lys Gln Leu Lys Leu Leu Phe Leu Ser Arg Asn His Leu Ser Ser Ile  
 180 185 190  
 Pro Ser Gly Leu Pro His Thr Leu Glu Glu Leu Arg Leu Asp Asp Asn  
 195 200 205  
 Arg Ile Ser Thr Ile Pro Leu His Ala Phe Lys Gly Leu Asn Ser Leu  
 210 215 220  
 Arg Arg Leu Val Leu Asp Gly Asn Leu Leu Ala Asn Gln Arg Ile Ala  
 225 230 235 240  
 Asp Asp Thr Phe Ser Arg Leu Gln Asn Leu Thr Glu Leu Ser Leu Val  
 245 250 255  
 Arg Asn Ser Leu Ala Ala Pro Pro Leu Tyr Leu Gln Asp Asn Ala Ile  
 260 265 270  
 Ser His Ile Pro Tyr Asn Thr Leu Ala Lys Met Arg Glu Leu Glu Arg  
 275 280 285  
 Leu Asp Leu Ser Asn Asn Asn Leu Thr Thr Leu Pro Arg Gly Leu Phe  
 290 295 300  
 Asp Asp Leu Gly Asn Leu Ala Gln Leu Leu Leu Arg Asn Asn Pro Trp  
 305 310 315 320  
 Phe Cys Gly Cys Asn Leu Met Trp Leu Arg Asp Trp Val Lys Ala Arg  
 325 330 335  
 Ala Ala Val Val Asn Val Arg Gly Leu Met Cys Gln Gly Pro Glu Lys  
 340 345 350  
 Val Arg Gly Met Ala Ile Lys Asp Ile Thr Ser Glu Val Glu Ser Val  
 355 360 365  
 Leu Arg Arg Ala Pro Gln Gly Gly Val Ala Asn Ala Ala Ala Lys Thr  
 370 375 380  
 Thr Ala Ser Asn His Ala Ser Ala Thr Thr Pro Gln Gly Ser Leu Phe  
 385 390 395 400  
 Thr Leu Lys Ala Lys Arg Pro Gly Leu Arg Leu Pro Asp Ser Asn Ile  
 405 410 415  
 Asp Tyr Pro Met Ala Thr Gly Asp Gly Ala Lys Thr Leu Ala Ile His  
 420 425 430

Val Lys Ala Leu Thr Ala Asp Ser Ile Arg Ile Thr Trp Lys Ala Thr  
 435 440 445  
 Leu Pro Ala Ser Ser Phe Arg Leu Ser Trp Leu Arg Leu Gly His Ser  
 450 455 460  
 Pro Ala Val Gly Ser Ile Thr Glu Thr Leu Val Gln Gly Asp Lys Thr  
 465 470 475 480  
 Glu Tyr Leu Leu Thr Ala Leu Glu Pro Lys Ser Thr Tyr Ile Ile Cys  
 485 490 495  
 Met Val Thr Met Glu Thr Ser Asn Ala Tyr Val Ala Asp Glu Thr Pro  
 500 505 510  
 Val Cys Ala Lys Ala Glu Thr Ala Asp Ser Tyr Gly Pro Thr Thr Thr  
 515 520 525  
 Leu Asn Gln Glu Gln Asn Ala Gly Pro Met Ala Ser Leu Pro Leu Ala  
 530 535 540  
 Gly Ile Ile Gly Gly Ala Val Ala Leu Val Phe Leu Phe Leu Val Leu  
 545 550 555 560  
 Gly Ala Ile Cys Trp Tyr Val His Gln Ala Gly Glu Leu Leu Thr Arg  
 565 570 575  
 Glu Arg Ala Tyr Asn Arg Gly Ser Arg Lys Lys Asp Asp Tyr Met Glu  
 580 585 590  
 Ser Gly Thr Lys Lys Asp Asn Ser Ile Leu Glu Ile Arg Gly Pro Gly  
 595 600 605  
 Leu Gln Met Leu Pro Ile Asn Pro Tyr Arg Ala Lys Glu Glu Tyr Val  
 610 615 620  
 Val His Thr Ile Phe Pro Ser Asn Gly Ser Ser Leu Cys Lys Ala Thr  
 625 630 635 640  
 His Thr Ile Gly Tyr Gly Thr Thr Arg Gly Tyr Arg Asp Gly Gly Ile  
 645 650 655  
 Pro Asp Ile Asp Tyr Ser Tyr Thr  
 660

<210> 69  
 <211> 26  
 <212> DNA  
 <213> Homo sapiens

<400> 69  
 caacgtgcag gtcattctacc tatacgt

26

<210> 70  
 <211> 25

<212> DNA  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:oligonucleotide  
primer

<400> 70  
gcccgtctca aaacactctc catct

25

<210> 71  
<211> 54  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:oligonucleotide  
primer

<400> 71  
Asn Pro Phe Asn Cys Asp Cys Glu Leu Arg Trp Leu Leu Arg Trp Leu  
1 5 10 15  
Arg Glu Thr Asn Pro Arg Arg Leu Glu Asp Gln Glu Asp Leu Arg Cys  
20 25 30  
Ala Ser Pro Glu Ser Leu Arg Gly Gln Pro Leu Leu Glu Leu Leu Pro  
35 40 45  
Ser Asp Phe Ser Cys Pro  
50

<210> 72  
<211> 84  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:consensus  
sequence

<400> 72  
Pro Ser Ala Pro Thr Asn Leu Thr Val Thr Asp Val Thr Ser Thr Ser  
1 5 10 15  
Leu Thr Leu Ser Trp Ser Pro Pro Thr Gly Asn Gly Pro Ile Thr Gly  
20 25 30  
Tyr Glu Val Thr Tyr Arg Gln Pro Lys Asn Gly Gly Glu Trp Asn Glu  
35 40 45  
Leu Thr Val Pro Gly Thr Thr Thr Ser Tyr Thr Leu Thr Gly Leu Lys  
50 55 60  
Pro Gly Thr Glu Tyr Glu Val Arg Val Gln Ala Val Asn Gly Gly Gly



145                      150                      155                      160  
 Gly Cys Cys Cys Cys Thr Cys Gly Gly Thr Gly Thr Gly Cys Cys Gly  
    165                      170                      175  
 Cys Thr Gly Cys Gly Ala Cys Ala Ala Cys Gly Gly Cys Thr Thr Cys  
    180                      185                      190  
 Ala Thr Cys Thr Ala Cys Thr Gly Cys Ala Ala Cys Gly Ala Cys Cys  
    195                      200                      205  
 Gly Gly Gly Gly Ala Cys Thr Cys Ala Cys Ala Thr Cys Cys Ala Thr  
    210                      215                      220  
 Cys Cys Cys Cys Gly Cys Ala Gly Ala Thr Ala Thr Cys Cys Cys Thr  
 225     230                      235                      240  
 Gly Ala Thr Gly Ala Cys Gly Cys Cys Ala Cys Cys Ala Cys Cys Cys  
    245                      250                      255  
 Thr Cys Thr Ala Thr Cys Thr Gly Cys Ala Gly Ala Ala Cys Ala Ala  
    260                      265                      270  
 Cys Cys Ala Gly Ala Thr Cys Ala Ala Cys Ala Ala Cys Gly Cys Thr  
    275                      280                      285  
 Gly Gly Cys Ala Thr Cys Cys Cys Cys Cys Ala Gly Gly Ala Cys Cys  
    290                      295                      300  
 Thr Cys Ala Ala Gly Ala Cys Cys Ala Ala Gly Gly Thr Cys Ala Ala  
 305     310                      315                      320  
 Cys Gly Thr Gly Cys Ala Gly Gly Thr Cys Ala Thr Cys Thr Ala Cys  
    325                      330                      335  
 Cys Thr Ala Thr Ala Cys Gly Ala Gly Ala Ala Thr Gly Ala Cys Cys  
    340                      345                      350  
 Thr Gly Gly Ala Thr Gly Ala Gly Thr Thr Cys Cys Cys Cys Ala Thr  
    355                      360                      365  
 Cys Ala Ala Cys Cys Thr Gly Cys Cys Cys Cys Gly Cys Thr Cys Cys  
    370                      375                      380  
 Cys Thr Cys Cys Gly Gly Gly Ala Gly Cys Thr Gly Cys Ala Cys Cys  
 385     390                      395                      400  
 Thr Gly Cys Ala Gly Gly Ala Cys Ala Ala Cys Ala Ala Thr Gly Thr  
    405                      410                      415  
 Gly Cys Gly Cys Ala Cys Cys Ala Thr Thr Gly Cys Cys Ala Gly Gly  
    420                      425                      430  
 Gly Ala Cys Thr Cys Gly Cys Thr Gly Gly Cys Cys Cys Gly Cys Ala  
    435                      440                      445  
 Thr Cys Cys Cys Gly Cys Thr Gly Cys Thr Gly Gly Ala Gly Ala Ala

450		455		460
Gly Cys Thr Gly Cys	Ala Cys Cys Thr Gly Gly	Ala Thr Gly Ala Cys		
465	470	475	480	
Ala Ala Cys Thr Cys	Cys Gly Thr Gly Thr Cys Cys	Ala Cys Cys Gly		
	485	490	495	
Thr Cys Ala Gly Cys	Ala Thr Thr Gly Ala Gly Gly	Ala Gly Gly Ala		
	500	505	510	
Cys Gly Cys Cys Thr	Thr Cys Gly Cys Cys Gly	Ala Cys Ala Gly Cys		
	515	520	525	
Ala Ala Ala Cys Ala	Gly Cys Thr Cys Ala Ala	Gly Cys Thr Gly Cys		
	530	535	540	
Thr Cys Thr Thr Cys	Cys Thr Gly Ala Gly Cys Cys	Gly Gly Ala Ala		
	545	550	555	560
Cys Cys Ala Cys Cys	Thr Gly Ala Gly Cys Ala	Gly Cys Ala Thr Cys		
	565	570	575	
Cys Cys Cys Thr Cys	Gly Gly Gly Gly Cys Thr	Gly Cys Cys Gly Cys		
	580	585	590	
Ala Cys Ala Cys Gly	Cys Thr Gly Gly Ala Gly	Gly Ala Gly Cys Thr		
	595	600	605	
Gly Cys Gly Gly Cys	Thr Gly Gly Ala Thr Gly	Ala Cys Ala Ala Cys		
	610	615	620	
Cys Gly Cys Ala Thr	Cys Thr Cys Cys Ala Cys	Cys Cys Ala Thr Cys	Cys	
	625	630	635	640
Cys Gly Cys Thr Gly	Cys Ala Thr Gly Cys	Cys Thr Thr Cys Ala	Ala	
	645	650	655	
Gly Gly Gly Cys Cys	Thr Cys Ala Ala Cys	Ala Gly Cys Cys Thr	Gly	
	660	665	670	
Cys Gly Gly Cys Gly	Cys Cys Thr Gly Gly	Thr Gly Cys Thr Gly	Gly	
	675	680	685	
Ala Cys Gly Gly Thr	Ala Ala Cys Cys Thr	Gly Cys Thr Gly Gly	Cys	
	690	695	700	
Cys Ala Ala Cys Cys	Ala Gly Cys Gly Cys	Ala Thr Cys Gly Cys	Cys	
	705	710	715	720
Gly Ala Cys Gly Ala	Cys Ala Cys Cys Thr	Thr Cys Ala Gly Cys	Cys	
	725	730	735	
Gly Cys Cys Thr Ala	Cys Ala Gly Ala Ala	Cys Cys Thr Cys Ala	Cys	
	740	745	750	
Ala Gly Ala Gly Cys	Thr Cys Thr Cys Gly	Cys Thr Gly Gly Thr	Gly	



755					760					765					
Cys	Gly	Cys	Ala	Ala	Thr	Thr	Cys	Gly	Cys	Thr	Gly	Gly	Cys	Cys	Gly
770						775					780				
Cys	Gly	Cys	Cys	Ala	Cys	Cys	Cys	Cys	Thr	Cys	Ala	Ala	Cys	Cys	Thr
785					790					795					800
Gly	Cys	Cys	Cys	Ala	Gly	Cys	Gly	Cys	Cys	Cys	Ala	Cys	Cys	Thr	Gly
				805					810					815	
Cys	Ala	Gly	Ala	Ala	Ala	Cys	Thr	Cys	Thr	Ala	Cys	Cys	Thr	Gly	Cys
			820					825					830		
Ala	Gly	Gly	Ala	Cys	Ala	Ala	Thr	Gly	Cys	Cys	Ala	Thr	Cys	Ala	Gly
			835				840					845			
Cys	Cys	Ala	Cys	Ala	Thr	Cys	Cys	Cys	Cys	Thr	Ala	Cys	Ala	Ala	Cys
	850					855					860				
Ala	Cys	Gly	Cys	Thr	Gly	Gly	Cys	Cys	Ala	Ala	Gly	Ala	Thr	Gly	Cys
865					870					875					880
Gly	Thr	Gly	Ala	Gly	Cys	Thr	Gly	Gly	Ala	Gly	Cys	Gly	Gly	Cys	Thr
				885					890					895	
Gly	Gly	Ala	Cys	Cys	Thr	Gly	Thr	Cys	Cys	Ala	Ala	Cys	Ala	Ala	Cys
			900					905					910		
Ala	Ala	Cys	Cys	Thr	Gly	Ala	Cys	Cys	Ala	Cys	Gly	Cys	Thr	Gly	Cys
			915				920					925			
Cys	Cys	Cys	Gly	Cys	Gly	Gly	Cys	Cys	Thr	Gly	Thr	Thr	Cys	Gly	Ala
	930					935					940				
Cys	Gly	Ala	Cys	Cys											
945															

<210> 75

<211> 674

<212> PRT

<213> Homo sapiens

<400> 75

Met	Val	Val	Ala	His	Pro	Thr	Ala	Thr	Ala	Thr	Thr	Thr	Pro	Thr	Ala
1				5					10					15	

Thr	Val	Thr	Ala	Thr	Val	Val	Met	Thr	Thr	Ala	Thr	Met	Asp	Leu	Arg
			20					25					30		

Asp	Trp	Leu	Phe	Leu	Cys	Tyr	Gly	Leu	Ile	Ala	Phe	Leu	Thr	Glu	Val
		35					40					45			

Ile	Asp	Ser	Thr	Thr	Cys	Pro	Ser	Val	Cys	Arg	Cys	Asp	Asn	Gly	Phe
	50					55					60				

Ile	Tyr	Cys	Asn	Asp	Arg	Gly	Leu	Thr	Ser	Ile	Pro	Ala	Asp	Ile	Pro	65	70	75	80
Asp	Asp	Ala	Thr	Thr	Leu	Tyr	Leu	Gln	Asn	Asn	Gln	Ile	Asn	Asn	Ala	85	90	95	
Gly	Ile	Pro	Gln	Asp	Leu	Lys	Thr	Lys	Val	Asn	Val	Gln	Val	Ile	Tyr	100	105	110	
Leu	Tyr	Glu	Asn	Asp	Leu	Asp	Glu	Phe	Pro	Ile	Asn	Leu	Pro	Arg	Ser	115	120	125	
Leu	Arg	Glu	Leu	His	Leu	Gln	Asp	Asn	Asn	Val	Arg	Thr	Ile	Ala	Arg	130	135	140	
Asp	Ser	Leu	Ala	Arg	Ile	Pro	Leu	Leu	Glu	Lys	Leu	His	Leu	Asp	Asp	145	150	155	160
Asn	Ser	Val	Ser	Thr	Val	Ser	Ile	Glu	Glu	Asp	Ala	Phe	Ala	Asp	Ser	165	170	175	
Lys	Gln	Leu	Lys	Leu	Leu	Phe	Leu	Ser	Arg	Asn	His	Leu	Ser	Ser	Ile	180	185	190	
Pro	Ser	Gly	Leu	Pro	His	Thr	Leu	Glu	Glu	Leu	Arg	Leu	Asp	Asp	Asn	195	200	205	
Arg	Ile	Ser	Thr	Ile	Pro	Leu	His	Ala	Phe	Lys	Gly	Leu	Asn	Ser	Leu	210	215	220	
Arg	Arg	Leu	Val	Leu	Asp	Gly	Asn	Leu	Leu	Ala	Asn	Gln	Arg	Ile	Ala	225	230	235	240
Asp	Asp	Thr	Phe	Ser	Arg	Leu	Gln	Asn	Leu	Thr	Glu	Leu	Ser	Leu	Val	245	250	255	
Arg	Asn	Ser	Leu	Ala	Ala	Pro	Pro	Leu	Asn	Leu	Pro	Ser	Ala	His	Leu	260	265	270	
Gln	Lys	Leu	Tyr	Leu	Gln	Asp	Asn	Ala	Ile	Ser	His	Ile	Pro	Tyr	Asn	275	280	285	
Thr	Leu	Ala	Lys	Met	Arg	Glu	Leu	Glu	Arg	Leu	Asp	Leu	Ser	Asn	Asn	290	295	300	
Asn	Leu	Thr	Thr	Leu	Pro	Arg	Gly	Leu	Phe	Asp	Asp	Leu	Gly	Asn	Leu	305	310	315	320
Ala	Gln	Leu	Leu	Leu	Arg	Asn	Asn	Pro	Trp	Phe	Cys	Gly	Cys	Asn	Leu	325	330	335	
Met	Trp	Leu	Arg	Asp	Trp	Val	Lys	Ala	Arg	Ala	Ala	Val	Val	Asn	Val	340	345	350	
Arg	Gly	Leu	Met	Cys	Gln	Gly	Pro	Glu	Lys	Val	Arg	Gly	Met	Ala	Ile	355	360	365	

Lys Asp Ile Thr Ser Glu Met Asp Glu Cys Phe Glu Thr Gly Pro Gln  
 370 375 380  
 Gly Gly Val Ala Asn Ala Ala Ala Lys Thr Thr Ala Ser Asn His Ala  
 385 390 395 400  
 Ser Ala Thr Thr Pro Gln Gly Ser Leu Phe Thr Leu Lys Ala Lys Arg  
 405 410 415  
 Pro Gly Leu Arg Leu Pro Asp Ser Asn Ile Asp Tyr Pro Met Ala Thr  
 420 425 430  
 Gly Asp Gly Ala Lys Thr Leu Ala Ile His Val Lys Ala Leu Thr Ala  
 435 440 445  
 Asp Ser Ile Arg Ile Thr Trp Lys Ala Thr Leu Pro Ala Ser Ser Phe  
 450 455 460  
 Arg Leu Ser Trp Leu Arg Leu Gly His Ser Pro Ala Val Gly Ser Ile  
 465 470 475 480  
 Thr Glu Thr Leu Val Gln Gly Asp Lys Thr Glu Tyr Leu Leu Thr Ala  
 485 490 495  
 Leu Glu Pro Lys Ser Thr Tyr Ile Ile Cys Met Val Thr Met Glu Thr  
 500 505 510  
 Ser Asn Ala Tyr Val Ala Asp Glu Thr Pro Val Cys Ala Lys Ala Glu  
 515 520 525  
 Thr Ala Asp Ser Tyr Gly Pro Thr Thr Thr Leu Asn Gln Glu Gln Asn  
 530 535 540  
 Ala Gly Pro Met Ala Ser Leu Pro Leu Ala Gly Ile Ile Gly Gly Ala  
 545 550 555 560  
 Val Ala Leu Val Phe Leu Phe Leu Val Leu Gly Ala Ile Cys Trp Tyr  
 565 570 575  
 Val His Gln Ala Gly Glu Leu Leu Thr Arg Glu Arg Ala Tyr Asn Arg  
 580 585 590  
 Gly Ser Arg Glu Lys Asp Asp Tyr Met Glu Ser Gly Thr Lys Lys Asp  
 595 600 605  
 Asn Ser Ile Leu Glu Ile Arg Gly Pro Gly Leu Gln Met Leu Pro Ile  
 610 615 620  
 Asn Pro Tyr Arg Ala Lys Glu Glu Tyr Val Val His Thr Ile Phe Pro  
 625 630 635 640  
 Ser Asn Gly Ser Ser Leu Cys Lys Ala Thr His Thr Ile Gly Tyr Gly  
 645 650 655  
 Thr Thr Arg Gly Tyr Arg Asp Gly Gly Ile Pro Asp Ile Asp Tyr Ser  
 660 665 670

Tyr Thr

<210> 76  
<211> 31  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:consensus  
sequence

<400> 76  
Ala Cys Pro Arg Glu Cys Thr Cys Ser Pro Phe Gly Leu Val Val Asp  
1 5 10 15  
Cys Ser Gly Arg Gly Leu Thr Leu Glu Val Pro Arg Asp Leu Pro  
20 25 30

<210> 77  
<211> 23  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:consensus  
sequence

<400> 77  
Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro  
1 5 10 15  
Gly Leu Phe Ser Asn Leu Pro  
20

<210> 78  
<211> 23  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:consensus  
sequence

<400> 78  
Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro  
1 5 10 15  
Gly Leu Phe Ser Asn Leu Pro  
20

<210> 79  
<211> 23

<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:consensus  
sequence

<400> 79  
Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro  
1 5 10 15

Gly Leu Phe Ser Asn Leu Pro  
20

<210> 80  
<211> 23  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:consensus  
sequence

<400> 80  
Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro  
1 5 10 15

Gly Leu Phe Ser Asn Leu Pro  
20

<210> 81  
<211> 23  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:consensus  
sequence

<400> 81  
Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro  
1 5 10 15

Gly Leu Phe Ser Asn Leu Pro  
20

<210> 82  
<211> 23  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:consensus  
sequence

<400> 82  
Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro  
1 5 10 15

Gly Leu Phe Ser Asn Leu Pro  
20

<210> 83  
<211> 23  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:consensus  
sequence

<400> 83  
Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro  
1 5 10 15

Gly Leu Phe Ser Asn Leu Pro  
20

<210> 84  
<211> 23  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence:consensus  
sequence

<400> 84  
Asn Leu Glu Glu Leu Asp Leu Ser Asn Asn Leu Thr Ser Leu Pro Pro  
1 5 10 15

Gly Leu Phe Ser Asn Leu Pro  
20